```
            |||||  |||||||||||||||||||  ||  ||  ||||||||||||  ||||||||||||||||||||
Db    2101  TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160


Qy    2206  AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2265
            |||||  |||||  ||||||||||||||||||||||||||  |||||  ||||||||||||  |||||
Db    2161  AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220


Qy    2266  CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2325
            ||||  |||   ||||||||||||||||||||||||||||  |||  |  |||||||  ||  ||||||
Db    2221  CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280


Qy    2326  GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2385
            ||||||||||  |||||||||||||||||||||||||||  ||  |||||||||||||  |||||||||
Db    2281  GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340


Qy    2386  AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2445
            ||||||||  |||||||||||||||||||||||||||||||  ||  |||||||||||||||||||  |||||||
Db    2341  AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400


Qy    2446  CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2505
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  |||||  ||
Db    2401  CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAGATCATC 2460


Qy    2506  TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2565
             .    ||||  |||||||||||||  ||  |||||  |||||||||  |  |||||  ||||||||||||
Db    2461  GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520


Qy    2566  CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2625
            ||||||||||||||||||  ||||||||||||||||||||||||||||||||||  ||||||||||||||
Db    2521  CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580


Qy    2626  AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2685
            |||||  |||||||||  |||||||||||||||||||||||||||||  |||||||||||  |||||  |||
Db    2581  AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640


Qy    2686  GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGC 2745
            ||  |||||||||||  |||||  ||||||||||||||||||  ||||||||||  |||||  ||||  |
Db    2641  GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGAGAC 2700


Qy    2746  CGGCCAG 2752
            |  |||||
Db    2701  CAGCCAG 2707
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RESULT 2
US-09-306-902A-1
; Sequence 1, Application US/09306902A
; Patent No. 6277585
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay
;   Masu, Masayuki
;   Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 9
;   CORRESPONDENCE ADDRESS:

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  Query Match              82.1%;  Score 2259;  DB 3;  Length 3014;
  Best Local Similarity    89.7%;  Pred. No. 0;
  Matches 2427;  Conservative   0;  Mismatches  280;  Indels   0;  Gaps   0;


Qy        46 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 105
             ||||||||||||||||||||||||||||||| ||||||||||||||||||||||||| |||||
Db         1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGTGCTCCTGGGCATAGTCCTCGCCGCCTGGCTT 60


Qy       106 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 165
             || || ||||||||||||||||||||||||| ||||||||| ||||||||| |||||||||||
Db        61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120


Qy       166 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 225
             |||||||| .||||||||||||||| ||||| ||||| ||||||||| ||||||||||||||
Db       121 GACCTGCTGCCCCACTTCCTGGTAGAGCCTGAGGACGTGTACATTGTCAAGAACAAGCCG 180


Qy       226 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 285
             ||| || | ||||||||||| ||||| ||||| |||||||||||||||||||| |||||
Db       181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240


Qy       286 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 345
             ||||| ||||||||| || |||||| || || ||||||||| ||| | |||||| || |||||
Db       241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300


Qy       346 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 405
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              ||||||||||||||||  ||  ||  ||  ||  ||||||||||||  |||||  |||||  |||||||||||
Db      301   ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG   360

Qy      406   GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC   465
              ||||||||||||||||||||  ||||||||||||||||||||||||  ||||||||  |||||||||||||||
Db      361   GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC   420

Qy      466   TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG   525
              ||||||||  ||  |||     ||||||||||||||  ||||||||||||  ||||||||||||  |||
Db      421   TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG   480

Qy      526   TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG   585
              ||  |||||||||  |||||  ||  ||  ||  ||  ||  ||  ||  ||  |||||  |||||  |||
Db      481   TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCCAGAAGGAATCCCCCCAGCTGAG   540

Qy      586   GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC   645
              ||||||||||  ||  ||  ||||||||||  ||||||||  |||||  ||  |||||||||  ||||||
Db      541   GTGGAGTGGCTTCGAAATGAGGACCTCGTGGACCCCTCCCTCGATCCCAATGTGTACATC   600

Qy      646   ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC   705
              ||||||||||||||||  ||  ||||||  ||||||||||  ||  ||||||||||||||||||||||
Db      601   ACGCGGGAGCACAGCCTAGTCGTGCGTCAGGCCCGCCTGGCCGACACGGCCAACTACACC   660

Qy      706   TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC   765
              ||  ||||||||||||||||||  ||  |||||||||  |||  ||||  ||  ||  |||||  ||  ||
Db      661   TGTGTGGCCAAGAACATCGTAGCCCGTCGCCGAAGCACCTCTGCAGCGGTCATTGTTTAT   720

Qy      766   GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC   825
              ||||||||||||||||||||||||||||  ||||||||||||||||||||||||||||||
Db      721   GTGAACGGTGGGTGGTCGACGTGGACTGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGT   780

Qy      826   GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC   885
              |||||||||||||||||||||||||||||||||||||||||||||  ||||||||||||||||||  |||
Db      781   GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC   840

Qy      886   TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC   945
              ||||||||||||||||||||||||||||||||||||||||||||  ||||||||||  ||  ||  |||
Db      841   TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACTCTGTGCCCAGTGGATGGGAGC   900

Qy      946   TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT   1005
              |||||     |||||||  ||||||||  ||||||||||||||  ||||||||||||||||||||||||||
Db      901   TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC   960

Qy      1006  GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC   1065
              |||||||||||||||||||||||||||  |||||  |||||||||  |  |||     ||||||||||||
Db      961   GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC   1020

Qy      1066  ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC   1125
              ||||||||||||||||||||||||||||  |  ||||     |||||||  ||||  |||||||||||
Db      1021  ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT   1080

Qy      1126  CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC   1185
              |||||  |  ||||||     ||||  |||||  ||  |||||     ||  ||||||  ||||  |  |||
Db      1081  CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA   1140

Qy      1186  CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC   1245
              |||  ||||  ||  ||  ||||||||  ||||||||||  |||||||||  |||||||||||  |||
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Db   1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC 1200

Qy   1246 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1305
          ||||| |||||||||||| |||||||||||||||||||||||||||||||||| ||||||
Db   1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC 1260

Qy   1306 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1365
          |||||||||| ||||||||||||||||||||||||| |||||||||||||||| ||| | ||
Db   1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG 1320

Qy   1366 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1425
          |||||||| |||||||||||||||||||||| | |||||| ||||||||||||||| |||||
Db   1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG 1380

Qy   1426 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1485
          | |||||||| || ||||||||||||||||| ||||||||||||| ||||| ||||||||
Db   1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC 1440

Qy   1486 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1545
          |||||||||||||| ||||||||| || ||||||||||||| |||||||||||||| ||||
Db   1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC 1500

Qy   1546 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1605
          ||||||||||||||||||||||||||||||||||||||||||||||| || |||||||||||
Db   1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC 1560

Qy   1606 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1665
          || ||||| ||||||||| |||||||| ||||||| |||||| |||||||||| |||||||
Db   1561 ATACCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG 1620

Qy   1666 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1725
          || |||||||||||||||||||||||||||||||||||||||||||||||| |||||||||
Db   1621 CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC 1680

Qy   1726 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1785
          ||||| ||||| || |||||||||||||||||||||||||| || ||||||||||||||
Db   1681 TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA 1740

Qy   1786 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1845
          |||||||||||||||||||||||| || ||||||||||||||||| |||||||||||| |||
Db   1741 GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG 1800

Qy   1846 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1905
          |||||||||||||||| || |||||| | || |||||||||||||||||||||||||||||
Db   1801 GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG 1860

Qy   1906 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1965
          ||| | ||||||||| ||||||||||||||||||||||||||||||||||||| |||||||
Db   1861 GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC 1920

Qy   1966 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 2025
          |||||||||||||||| ||||||||||| || |||||||||||| || ||||||| ||
Db   1921 CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG 1980

Qy   2026 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2085
          ||||| |||||||||||||| || ||||||| || ||||||||| || |||||||||
Db   1981 TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG 2040
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Qy    2086 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2145
           ||||||||||||||||||||||||||| || ||||||||||||||||||||| || |||||||||
Db    2041 GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC 2100

Qy    2146 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2205
           ||||| ||||||||||||||||||| || || ||||||||||| |||||||||||||||||||
Db    2101 TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160

Qy    2206 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2265
           ||||| ||||| |||||||||||||||||||||||| ||||| |||||||||| |||||
Db    2161 AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy    2266 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2325
           |||| ||| |||||||||||||||||||||||||| ||| | |||||||| || ||||||
Db    2221 CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280

Qy    2326 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2385
           |||||||||| ||||||||||||||||||||||||| || ||||||||||||| |||||||||
Db    2281 GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy    2386 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2445
           ||||||||| ||||||||||||||||||||  || |||||||||||||||||| |||||||
Db    2341 AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy    2446 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2505
           |||||||||||||||||||||||||||||||||||||||||||||||||||| |||||. ||
Db    2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy    2506 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2565
           |||| |||||||||||||||| || ||||| ||||||||| | ||||| |||||||||||
Db    2461 GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy    2566 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2625
           |||||||||||||||||| |||||||||||||||||||||||||||| |||||||||||||||
Db    2521 CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy    2626 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2685
           ||||| ||||||||| ||||||||||||||||||||||||| ||||||||||| ||||| |||
Db    2581 AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy    2686 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGC 2745
           || ||||||||||| ||||| ||||||||||||||| |||||||||||| ||||| |||| |
Db    2641 GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGAGAC 2700

Qy    2746 CGGCCAG 2752
           | |||||
Db    2701 CAGCCAG 2707
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RESULT 3
US-08-808-982-2
; Sequence 2, Application US/08808982
; Patent No. 5939271
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   APPLICANT: Leonardo, E. David

```
  Query Match            56.8%;  Score 1562.4;  DB 2;  Length 1787;
  Best Local Similarity  98.5%;  Pred. No. 0;
  Matches 1661; Conservative    0;  Mismatches  16;  Indels   9;  Gaps    8;

Qy     1070 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1129
            |||||||||||||||||||||||| ||||||| ||||||||||||||||||||||||||||
Db        1 GCAACTGTACCAGTGACCTCTG-GTACACACTGCTTCTGGCCCTGAGGACGTGGCCCTCT 59

Qy     1130 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 1189
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119

Qy     1190 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1249
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      120 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 179

Qy     1250 CAGGCTTCCAGCCCGTCAGCATC-AAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 1308
            ||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Db      180 CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239
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Qy   1309 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1368
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    240 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299

Qy   1369 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1428
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    300 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359

Qy   1429 GGCCGCCACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1488
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    360 GGCCGCCACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419

Qy   1489 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 1548
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    420 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 479

Qy   1549 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1608
          |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db    480 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATC 539

Qy   1609 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1668
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    540 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599

Qy   1669 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1728
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    600 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy   1729 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1788
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    660 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy   1789 CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1848
          ||||||||||||||||||||||||||  |||||||||||||||||||||||||| |||||||||
Db    720 CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy   1849 GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1908
          |||||  |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    779 GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy   1909 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1968
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    838 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy   1969 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 2028
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||.|.|
Db    898 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957

Qy   2029 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2088
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    958 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017

Qy   2089 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTC 2148
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1018 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

Qy   2149 AAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAG 2208
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               ||||||||||||||||||||||   |||        ||||||||||||||||||||||||||||||||
Db     1077   AAGGACAGTTACCACAACCT--GCCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy     2209   AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2268
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1135   AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy     2269   CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2328
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1195   CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy     2329   TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2388
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1255   TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 1314

Qy     2389   ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 2448
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1315   ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 1374

Qy     2449   GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 2508
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1375   GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 1434

Qy     2509   AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2568
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1435   AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 1494

Qy     2569   CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 2628
               ||||||||||||||||||||||||||||||||||||||||||||||||.|||||||||||
Db     1495   CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

Qy     2629   CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 2688
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1555   CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 1614

Qy     2689   GGACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGAGGCC 2746
               ||      |||||||   ||||  ||||  ||  |||||||| ||||||||||||||||||
Db     1615   GGGACTGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGAGGCC 1674

Qy     2747   GGCCAG 2752
               ||||||
Db     1675   GGCCAG 1680
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RESULT 4
US-09-306-902A-2
; Sequence 2, Application US/09306902A
; Patent No. 6277585
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay
;   Masu, Masayuki
;   Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 9
;   CORRESPONDENCE ADDRESS:

US-09-306-902A-2

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  Query Match              56.8%;  Score 1562.4;  DB 3;  Length 1787;
  Best Local Similarity    98.5%;  Pred. No. 0;
  Matches 1661;  Conservative    0;  Mismatches   16;  Indels   9;  Gaps    8;


Qy      1070 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1129
             ||||||||||||||||||||||||| ||||||| |||||||||||||||||||||||||||
Db         1 GCAACTGTACCAGTGACCTCTG-GTACACACTGCTTCTGGCCCTGAGGACGTGGCCCTCT 59


Qy      1130 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 1189
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119


Qy      1190 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1249
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       120 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 179


Qy      1250 CAGGCTTCCAGCCCGTCAGCATC-AAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 1308
             ||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Db       180 CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239


Qy      1309 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1368
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       240 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299


Qy      1369 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1428
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                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     300 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359

Qy    1429 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1488
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     360 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419

Qy    1489 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 1548
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     420 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 479

Qy    1549 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1608
                  |||||||||||||||||||||||||||||||||||||||||||| |||||||||||||||
Db     480 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATC 539

Qy    1609 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1668
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     540 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599

Qy    1669 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1728
                  ||||.||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     600 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy    1729 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1788
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     660 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy    1789 CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1848
                  ||||||||||||||||||||||||    |||||||||||||||||||||    ||||||||
Db     720 CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy    1849 GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1908
                  |||||  |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     779 GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy    1909 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1968
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     838 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy    1969 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 2028
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     898 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957

Qy    2029 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2088
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     958 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017

Qy    2089 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTC 2148
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1018 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

Qy    2149 AAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAG 2208
                  |||||||||||||||||||||   |||          |||||||||||||||||||||||
Db    1077 AAGGACAGTTACCACAACCT--GCCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy    2209 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2268
                  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db    1135 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy    2269 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2328
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1195 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy    2329 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2388
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1255 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 1314

Qy    2389 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 2448
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1315 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 1374

Qy    2449 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 2508
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1375 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 1434

Qy    2509 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2568
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1435 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 1494

Qy    2569 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 2628
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1495 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

Qy    2629 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 2688
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1555 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 1614

Qy    2689 GGACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGAGGCC 2746
           ||      |||||||   ||||  |||| || |||||||||  ||||||||||||||||||
Db    1615 GGGACTGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGAGGCC 1674

Qy    2747 GGCCAG 2752
           ||||||
Db    1675 GGCCAG 1680
```

RESULT 5
US-08-808-982-3
; Sequence 3, Application US/08808982
; Patent No. 5939271
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   APPLICANT: Leonardo, E. David
;   APPLICANT: Hink, Lindsay
;   APPLICANT: Masu, Masayuki
;   APPLICANT: Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 8
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;   STREET: 268 BUSH STREET, SUITE 3200
;   CITY: SAN FRANCISCO
;   STATE: CALIFORNIA
;   COUNTRY: USA

```
;        ZIP:  94104
;     COMPUTER READABLE FORM:
;       MEDIUM TYPE:  Floppy disk
;       COMPUTER:  IBM PC compatible
;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  PatentIn Release #1.0, Version #1.30
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/808,982
;       FILING DATE:
;       CLASSIFICATION:  530
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  OSMAN, RICHARD A
;       REGISTRATION NUMBER:  36,627
;       REFERENCE/DOCKET NUMBER:  UC96-217
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  (415) 343-4341
;       TELEFAX:  (415) 343-4342
;   INFORMATION FOR SEQ ID NO:  3:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  2831 base pairs
;       TYPE:  nucleic acid
;       STRANDEDNESS:  double
;       TOPOLOGY:  linear
;     MOLECULE TYPE:  cDNA
US-08-808-982-3
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   Query Match            30.6%;   Score 841.4;  DB 2;   Length 2831;
   Best Local Similarity  60.0%;   Pred. No. 1.3e-178;
   Matches 1638;  Conservative    0;  Mismatches  961;  Indels  130;  Gaps    9;

Qy        143 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 202
              || | | || || | |||| | ||| || |||||||| |||| || |||||||
Db        104 ACTCCTTCCCATCAGCACCCGCGGAGCAGCTGCCTCACTTCCTGCTGGAACCAGAGGATG 163

Qy        203 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 262
              ||||||| ||||||||||||||| |     ||| ||| | || ||||| ||||
Db        164 CCTACATCGTAAAGAACAAGCCAGTGGAATTGCACTGCCGAGCCTTCCCTGCCACACAGA 223

Qy        263 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 322
              ||| ||||||||| || || |||||||| ||||| | ||||||| | ||    |||
Db        224 TCTACTTCAAGTGTAATGGCGAGTGGGTTAGCCAGAAAGGCCACGTCACGCAGGAGAGCC 283

Qy        323 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 382
              || | | || || ||| |      ||||| | || | || || |||||||||
Db        284 TGGATGAGGCCACAGGCTTGCGAATACGAGAGGTGCAGATAGAGGTGTCGCGGCAGCAGG 343

Qy        383 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 442
              | ||| | | || ||||| ||||| ||||||||| |||||||||| ||||||||| ||||
Db        344 TGGAGGAACTTTTTGGGCTCGAGGACTACTGGTGTCAGTGCGTGGCCTGGAGCTCTTCGG 403

Qy        443 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 502
              | |||||||||||| |||||||||||||| |||    |||||||||||||||| || |
Db        404 GAACCACCAAGAGTCGCCGAGCCTACATCCGCATTGCCTACTTGCGCAAGAACTTTGACC 463

Qy        503 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 562
              ||||||| ||||| ||||||||| || |||| || |   || | |||| ||||| ||||
Db        464 AGGAGCCTCTGGCGAAGGAGGTACCCTTGGATCATGAGGTCCTTCTGCAGTGCCGCCCAC 523
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Qy     563 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 622
               | |||||  |  |||    || |||||||||  ||||||     ||| || ||  |   | ||  ||
Db     524 CAGAGGGAGTGCCTGTGGCTGAGGTGGAATGGCTCAAGAATGAAGATGTCATCGATCCCG 583

Qy     623 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 682
               | |  |||| | ||  |      |||| ||  ||||| |||  |  | || ||||| ||||
Db     584 CTCAGGACACTAACTTCCTGCTCACCATTGACCACAACCTCATCATCCGCCAGGCGCGCC 643

Qy     683 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 742
               |  |  ||||| ||||||||||||| ||||| ||||| ||  |||||     ||||| |||
Db     644 TCTCAGACACAGCCAACTACACCTGTGTGGCAAAGAATATTGTGGCCAAGCGCCGGAGCA 703

Qy     743 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 802
               |  |  ||  |  |||||||||||| |||||||||  | |||||| |   ||| | || |||||
Db     704 CGACGGCCACAGTCATCGTCTATGTGAACGGAGGTTGGTCCAGCTGGGCAGAATGGTCAC 763

Qy     803 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 862
               ||||       | ||||  || ||  || || ||||||||||| |    |||  |||||||||||| |
Db     764 CCTGCTCTAACCGCTGCGGCCGAGGTTGGCAGAAACGTACTAGGACCTGCACCAACCCAG 823

Qy     863 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 922
               | | || ||||| || || || || ||||| ||||| ||  |   | |||||| || || ||| |||
Db     824 CCCCACTCAATGGAGGTGCCTTCTGCGAGGGACAGGCTTGCCAGAAGACGGCTTGCACCA 883

Qy     923 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 982
               || ||||||||||| || ||    |||| |||||||||||||||||| |||||   |   |
Db     884 CCGTGTGCCCAGTGGATGGAGCGTGGACTGAGTGGAGCAAGTGGTCCGCCTGCAGCACAG 943

Qy     983 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 1042
               |  ||  |  |||||||| |||||  ||||| |||||  |  ||  |  | ||||  ||||||||   |
Db     944 AGTGTGCGCACTGGCGCAGCCGCGAGTGCATGGCACCGCCGCCCCAGAACGGAGGCCGTG 1003

Qy    1043 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1102
               |  |||      || ||     || |||  ||     |||||  ||  || || || | |  |  |
Db    1004 ACTGCAGCGGGACGCTACTTGACTCCAAGAACTGCACCGATGGGCTGTGCGTGCTGAATC 1063

Qy    1103 CTTCTGGCC-------------------------------CTGAGGACGTGGCCCTCT 1129
               |                                          |  |  |||||||| || |
Db    1064 AGAGAACTCTAAACGACCCTAAAAGCCGCCCCCTGGAGCCGTCGGGAGACGTGGCGCTGT 1123

Qy    1130 ATGTGGGCCTC---ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCC 1186
               ||| |||||||    |  |||||  ||    |||| |  ||  || ||  |||         |
Db    1124 ATGCGGGCCTCGTGGTGGCCGTCTTTGTGGTTCTGGCAGTTCTCATGGCTGTAGGAGTGA 1183

Qy    1187 TCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT---C 1243
               ||||  ||  | | || ||   | |   |  ||| | || |  |||||||| ||  |   |
Db    1184 TCGTGTACCGGAGAAACTGCCGGGACTTCGACACGGACATCACTGACTCCTCTGCTGCCC 1243

Qy    1244 TCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGC 1303
               ||||    || || |||| |||||| |||||| ||||||| |     |   |  |||||| || || |
Db    1244 TCACTGGTGGTTTCCACCCCGTCAACTTCAAGACTGCAAGGCCCAGCAACCCACAGCTCC 1303

Qy    1304 T------CACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCT 1357
               |        ||  ||  |||||||| |   ||| |   || |||||| ||   |  |  |
Db    1304 TGCACCCATCCGCCCCTCCGGACCTAACGGCCAGTGCTGGCATCTACCGCGGACCTGTGT 1363
```

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Qy    1358 GTCCCCGGCAGGA----------------------------------------------TG 1372
              | ||| ||||||
Db    1364 ATGCCCTGCAGGACTCTGCCGACAAGATCCCTATGACTAATTCACCCCTTCTGGATCCCT 1423

Qy    1373 GGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCC---------CCTGG 1423
              ||||||||| |||| || || ||    |       |    || || |         |||||
Db    1424 TGCCCAGCCTCAAGATCAAGGTCTATGACTCCAGCACCATCGGCTCTGGGGCTGGCCTGG 1483

Qy    1424 GTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCT 1483
              ||   ||   | ||   |||       |     | |||   |   |  |  || ||| |
Db    1484 CTGATGGAGCCGACCTGCTGGGTGTCTTACCACCCGGTACATACCCAGGCGATTTCTCCC 1543

Qy    1484 CCCGCCTCTCCACCCAGAAC---------------------------TACTTCCGCTCCC 1516
               | | | | || | ||                              || |||   ||
Db    1544 GGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCCTTGGTTCCCAGCACCTCCTGGGCC 1603

Qy    1517 TGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGGCTGA 1576
              | || |||| | ||||||| ||    | ||| |||||    | ||| || || |||||
Db    1604 TCCCTCGAGACCCCAGCAGCAGTGTCAGTGGCACCTTTGGTTGCCTGGGTGGGAGGCTGA 1663

Qy    1577 TGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAGGGAAGA 1636
               || ||     |||||  |||||||  |  | ||    |  ||||| |||| || |||
Db    1664 CCATTCCCGGCACAGGGGTCAGCCTGTTGGTACCAAATGGAGCCATTCCCCAGGGCAAGT 1723

Qy    1637 TCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCT 1696
              ||||||| | || ||      |  ||||| | ||| |    |       | ||| |
Db    1724 TCTATGACTTGTATCTACGTATCAACAAGACTGAAAGCACCCTCCCACTTTCGGAAGGTT 1783

Qy    1697 GTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGC 1756
              |||||  |  |||| |||   || | | ||| || ||| | ||| |||| ||   ||| |
Db    1784 CCCAGACAGTATTGAGCCCCTCGGTGACCTGCGGGCCCACGGGCCTCCTCCTGTGCCGCC 1843

Qy    1757 CAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCC 1816
              | ||  |||| || || |||||||| || ||    | | |||| ||||| | | | |
Db    1844 CTGTTGTCCTCACTGTGCCCCACTGTGCTGAAGTCATTGCCGGAGACTGGATCTTCCAGC 1903

Qy    1817 TCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGC 1876
              |||| |  ||| |     ||||| |||||||||| ||| ||    ||| |||||| | |
Db    1904 TCAAGACCCAGGCCCATCAGGGCCACTGGGAGGAGGTGGTGACTTTGGATGAGGAGACTC 1963

Qy    1877 CCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGC 1936
               |  || || |||||||||||||||| |||||| |    ||||| || || |    || ||||
Db    1964 TGAACACCCCCTGCTACTGCCAGCTAGAGGCTAAATCCTGCCACATCCTGTTGGACCAGC 2023

Qy    1937 TGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGC 1996
               .. ||||   || | |   ||| ||| || || | |    | ||| | |||||.||| ||| |||
Db    2024 TGGGTACCTACGTGTTCACGGGCGAGTCCTACTCCCGCTCCGCAGTCAAGCGGCTCCAGC 2083

Qy    1997 TGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTACTGCC 2056
              |       | || || || || |       |||||||||||||| |||||| |    || ||||||||||| |
Db    2084 TAGCCATCTTCGCCCCAGCCCTCTGCACCTCCCTGGAGTATAGTCTCAGGGTCTACTGTC 2143

Qy    2057 TGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGAC 2116
              || | |||||| | |   |||||| ||||||| || |||||||||| |     ||||| ||
Db    2144 TGGAGGACACTCCTGCAGCACTGAAGGAGGTCCTAGAGCTGGAGAGGACTCTGGGTGGCT 2203

Qy    2117 AGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGCGCCTAT 2176
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                     |  || |   |||||||      |      ||| ||| ||||||||||||||||||||||| || || |
Db      · 2204 ACTTGGTGGAGGAGCCCAAGACTTTGCTCTTTAAGGACAGTTACCACAACCTACG-CTCT 2262

Qy        2177 CCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGA 2236
                 || |||| ||  | |||     |||  |||| ||| || || || | ||  |||||||||
Db        2263 CCCTCCATGACATCCCCCATGCCCACTGGAGGAGCAAACTACTGGCCAAGTACCAGGAGA 2322

Qy      · 2237 TCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACCCTGG 2296
                |  ||||||  ||  ||  |  |  ||||| ||||   |||        | |||||||||| ||||||||||
Db        2323 TTCCCTTCTACCATGTGTGGAACGGCAGCCAGAAAGCCCTGCACTGCACTTTCACCCTGG 2382

Qy        2297 AGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGG 2356
                || |     ||||     | | | |||   | ||||  |||   | || ||| ||||||| || |
Db        2383 AGAGACATAGCCTAGCCTCCACTGAGTTCACCTGTAAGGTCTGCGTGCGGCAGGTAGAAG 2442

Qy        2357 GCGACGGGCAGAGCTTCAGCATCAACTTCAACATCAC---CAAGGACACAAGGTTTGCTG 2413
                |  || || ||||   |||     |  ||  || | |      | |   ||  |     |
Db        2443 GGGAAGGCCAGATTTTCCAGCTGCACACCACGCTGGCTGAGACGCCTGCTGGCTCCCTGG 2502

Qy        2414 AGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTCA 2473
                |    ||      |     ||     |  | | |  | | | |||| |||  |||||||||
Db        2503 ATGCACTCTGCTCTGCCCCTGGCAATGCTGCCACCACACAGCTGGGACCCTATGCCTTCA 2562

Qy        2474 AGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGCGGG 2533
                ||||  ||  |    ||| ||  ||||||||         |||||||||| | |||     ||||
Db        2563 AGATACCACTGTCCATCCGCCAGAAGATCTGCAACAGCCTGGACGCCCCCAACTCACGGG 2622

Qy        2534 GTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCTTTG 2593
                |     ||||||||||    |||| ||||| |||  | |||||| |  | || | || ||| |
Db        2623 GCAATGACTGGCGGCTGTTGGCACAGAAGCTCTCCATGGACCGGTACCTGAACTACTTCG 2682

Qy        2594 CCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCCCCA 2653
                || |||| | | || ||||||| | ||||| | ||||  | |||| |||||| || |||||
Db        2683 CCACCAAAGCTAGTCCCACAGGCGTGATCTTAGACCTCTGGGAAGCTCGGCAGCAGGATG 2742

Qy        2654 ACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTGGCC 2713
                | ||  ||||||| |   ||||| ||  |||   | | ||||| || ||    ||
Db        2743 ATGGGGACCTCAACAGCCTGGCCAGTGCCTTGGAGGAGATGGGCAAGAGTGAGATGCTGG 2802

Qy        2714 TCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
                |   ||       | || |  || ||||||
Db        2803 TAGCCATGACCACTGATGGCGATTGCTGA 2831


RESULT 6
US-09-306-902A-3
; Sequence 3, Application US/09306902A
; Patent No. 6277585
;    GENERAL INFORMATION:
;         APPLICANT: Tessier-Lavigne, Marc
;                    Leonardo, E. David
;                    Hink, Lindsay
;                    Masu, Masayuki
;                    Kazuko, Keino-Masu
;         TITLE OF INVENTION: Netrin Receptors
;         NUMBER OF SEQUENCES: 9
;         CORRESPONDENCE ADDRESS:
```

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   Query Match                30.6%;  Score 841.4;  DB 3;  Length 2831;
   Best Local Similarity      60.0%;  Pred. No. 1.3e-178;
   Matches 1638;  Conservative   0;  Mismatches  961;  Indels  130;  Gaps    9;

Qy        143 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 202
              || |  | ||    ||   | |||| | ||| || |||||||||| |||| || |||||||
Db        104 ACTCCTTCCCATCAGCACCCGCGGAGCAGCTGCCTCACTTCCTGCTGGAACCAGAGGATG 163

Qy        203 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 262
              |||||||| ||||||||||||||||| |   ||| ||| | || ||||| ||||
Db        164 CCTACATCGTAAAGAACAAGCCAGTGGAATTGCACTGCCGAGCCTTCCCTGCCACACAGA 223

Qy        263 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 322
              ||| ||||||||| || .|| |||||||| |||||   | ||||||| |  ||   |||
Db        224 TCTACTTCAAGTGTAATGGCGAGTGGGTTAGCCAGAAAGGCCACGTCACGCAGGAGAGCC 283

Qy        323 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 382
              || | |  || || ||| |     ||||| |  || | || || ||  |||||||||
Db        284 TGGATGAGGCCACAGGCTTGCGAATACGAGAGGTGCAGATAGAGGTGTCGCGGCAGCAGG 343

Qy        383 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 442
              | ||| |  | || |||||| ||||| ||||||||| |||||||||||| ||||||| ||||
Db        344 TGGAGGAACTTTTTGGGCTCGAGGACTACTGGTGTCAGTGCGTGGCCTGGAGCTCTTCGG 403

Qy        443 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 502
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                    |  |||||||||||||||        ||||||||||||||| |||      ||||||||||||||| || |
Db      404 GAACCACCAAGAGTCGCCGAGCCTACATCCGCATTGCCTACTTGCGCAAGAACTTTGACC 463


Qy      503 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 562
                ||||||| |||||  ||||||||  ||  |||| ||| |   ||| | |||| ||||| ||||
Db      464 AGGAGCCTCTGGCGAAGGAGGTACCCTTGGATCATGAGGTCCTTCTGCAGTGCCGCCCAC 523


Qy      563 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 622
                | ||||| | |||   || |||||||||| ||||||   ||| || ||  |   | || ||
Db      524 CAGAGGGAGTGCCTGTGGCTGAGGTGGAATGGCTCAAGAATGAAGATGTCATCGATCCCG 583


Qy      623 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 682
                | | |||| | || |     ||||    || ||||| |||   | | || |||||| ||||
Db      584 CTCAGGACACTAACTTCCTGCTCACCATTGACCACAACCTCATCATCCGCCAGGCGCGCC 643


Qy      683 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 742
                |  | ||||| |||||||||||||| ||||| ||||| || ||||| |       ||||| |||
Db      644 TCTCAGACACAGCCAACTACACCTGTGTGGCAAAGAATATTGTGGCCAAGCGCCGGAGCA 703


Qy      743 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 802
                | | || | |||||||||||| |||||||| || |||||| |   ||| | || |||||
Db      704 CGACGGCCACAGTCATCGTCTATGTGAACGGAGGTTGGTCCAGCTGGGCAGAATGGTCAC 763


Qy      803 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 862
                ||||     | ||||  ||  || || ||||||||||| |    ||| |||||||||||| |
Db      764 CCTGCTCTAACCGCTGCGGCCGAGGTTGGCAGAAACGTACTAGGACCTGCACCAACCCAG 823


Qy      863 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 922
                | || ||||| || || || |,| |||||  ||||| || |  | ||||| || || ||| |||
Db      824 CCCCACTCAATGGAGGTGCCTTCTGCGAGGGACAGGCTTGCCAGAAGACGGCTTGCACCA 883


Qy      923 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 982
                || |||||||||| || ||    ||||     ||||||||||||||| ||||| |     |
Db      884 CCGTGTGCCCAGTGGATGGAGCGTGGACTGAGTGGAGCAAGTGGTCCGCCTGCAGCACAG 943


Qy      983 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 1042
                | ||  | ||||||||| |||||||| ||||| |||||    | ||  | |||| |||||||| |
Db      944 AGTGTGCGCACTGGCGCAGCCGCGAGTGCATGGCACCGCCGCCCCAGAACGGAGGCCGTG 1003


Qy     1043 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1102
                | ||| || ||     || ||| || || |||||| ||| ||  || || || ||  | ||
Db     1004 ACTGCAGCGGGACGCTACTTGACTCCAAGAACTGCACCGATGGGCTGTGCGTGCTGAATC 1063


Qy     1103 CTTCTGGCC----------------------------------CTGAGGACGTGGCCCTCT 1129
                |                                        | |  ||||||||| || |
Db     1064 AGAGAACTCTAAACGACCCTAAAAGCCGCCCCCTGGAGCCGTCGGGAGACGTGGCGCTGT 1123


Qy     1130 ATGTGGGCCTC---ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCC 1186
                |||  |||||||     | | |||||   ||     |||| |  | | |  || || |||       |
Db     1124 ATGCGGGCCTCGTGGTGGCCGTCTTTGTGGTTCTGGCAGTTCTCATGGCTGTAGGAGTGA 1183


Qy     1187 TCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT---C 1243
                |||| ||  |  | | ||        ||   | ||| | ||  | |||||||||| ||   |   |
Db     1184 TCGTGTACCGGAGAAACTGCCGGGACTTCGACACGGACATCACTGACTCCTCTGCTGCCC 1243


Qy     1244 TCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGC 1303
                ||||    || ||||| ||||||| | ||||| |        |   |      |||||| || || |
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Db    1244 TCACTGGTGGTTTCCACCCCGTCAACTTCAAGACTGCAAGGCCCAGCAACCCACAGCTCC 1303

Qy    1304 T------CACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCT 1357
           |       ||  ||  |||||||| |   |||  |   || ||||| ||   | ||
Db    1304 TGCACCCATCCGCCCCTCCGGACCTAACGGCCAGTGCTGGCATCTACCGCGGACCTGTGT 1363

Qy    1358 GTCCCCGGCAGGA-----------------------------------------TG 1372
              | ||| ||||||
Db    1364 ATGCCCTGCAGGACTCTGCCGACAAGATCCCTATGACTAATTCACCCCTTCTGGATCCCT 1423

Qy    1373 GGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCC---------CCTGG 1423
           |||||||| |||| || || ||     |     |  || || |        |||||
Db    1424 TGCCCAGCCTCAAGATCAAGGTCTATGACTCCAGCACCATCGGCTCTGGGGCTGGCCTGG 1483

Qy    1424 GTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCT 1483
           || ||  | ||   |||     |     | |||   |  | |  | |  || ||| |
Db    1484 CTGATGGAGCCGACCTGCTGGGTGTCTTACCACCCGGTACATACCCAGGCGATTTCTCCC 1543

Qy    1484 CCCGCCTCTCCACCCAGAAC------------------------TACTTCCGCTCCC 1516
           | | | | || | ||                         || ||| ||
Db    1544 GGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCCTTGGTTCCCAGCACCTCCTGGGCC 1603

Qy    1517 TGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGGCTGA 1576
           | || |||| | ||||||| ||    | ||| |||||     | ||| || || ||||||
Db    1604 TCCCTCGAGACCCCAGCAGCAGTGTCAGTGGCACCTTTGGTTGCCTGGGTGGGAGGCTGA 1663

Qy    1577 TGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAGGGAAGA 1636
           || ||     ||||| ||||||| | | || | ||||| |||| |||| || ||
Db    1664 CCATTCCCGGCACAGGGGTCAGCCTGTTGGTACCAAATGGAGCCATTCCCCAGGGCAAGT 1723

Qy    1637 TCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCT 1696
           |||||||  |  || ||     | |||||  |  || |        |      |  || |
Db    1724 TCTATGACTTGTATCTACGTATCAACAAGACTGAAAGCACCCTCCCACTTTCGGAAGGTT 1783

Qy    1697 GTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGC 1756
           |||||  |  |||| |||  || || | ||| || ||| | || |||| || ||   ||| |
Db    1784 CCCAGACAGTATTGAGCCCCTCGGTGACCTGCGGGCCCACGGGCCTCCTCCTGTGCCGCC 1843

Qy    1757 CAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCC 1816
           | ||  ||||   || ||  || || |||||||| ||   ||  | |||||| | | | |
Db    1844 CTGTTGTCCTCACTGTGCCCCACTGTGCTGAAGTCATTGCCGGAGACTGGATCTTCCAGC 1903

Qy    1817 TCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGC 1876
           ||||| |  ||| |     ||||| ||||||||| ||| ||   ||| ||||||| | |
Db    1904 TCAAGACCCAGGCCCATCAGGGCCACTGGGAGGAGGTGGTGACTTTGGATGAGGAGACTC 1963

Qy    1877 CCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGC 1936
           | ||| || ||||||||||||||| |    ||||| || || |    || ||||
Db    1964 TGAACACCCCCTGCTACTGCCAGCTAGAGGCTAAATCCTGCCACATCCTGTTGGACCAGC 2023

Qy    1937 TGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGC 1996
           ||||    ||  |   |   ||| ||| ||  |   | || | |||||||| ||| |||
Db    2024 TGGGTACCTACGTGTTCACGGGCGAGTCCTACTCCCGCTCCGCAGTCAAGCGGCTCCAGC 2083

Qy    1997 TGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTACTGCC 2056
           |    | || || || ||   |||||||||||| |||||| |  || ||||||||||| |
Db    2084 TAGCCATCTTCGCCCCAGCCCTCTGCACCTCCCTGGAGTATAGTCTCAGGGTCTACTGTC 2143
```

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Qy    2057 TGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGAC 2116
           || | |||||| |  |   ||||| ||||||||| | ||||||||| |   ||||| ||
Db    2144 TGGAGGACACTCCTGCAGCACTGAAGGAGGTCCTAGAGCTGGAGAGGACTCTGGGTGGCT 2203

Qy    2117 AGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGCGCCTAT 2176
           |  || |  |||||||    |   ||| ||| |||||||||||||||||||||| || ||  |
Db    2204 ACTTGGTGGAGGAGCCCAAGACTTTGCTCTTTAAGGACAGTTACCACAACCTACG-CTCT 2262

Qy    2177 CCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGA 2236
           || |||| ||  | |||    ||| ||||  ||  ||  || || || || | || |||||||||
Db    2263 CCCTCCATGACATCCCCCATGCCCACTGGAGGAGCAAACTACTGGCCAAGTACCAGGAGA 2322

Qy    2237 TCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACCCTGG 2296
           | |||||| || ||  | ||||| ||||  ||     | ||||||||||| ||||||||||
Db    2323 TTCCCTTCTACCATGTGTGGAACGGCAGCCAGAAAGCCCTGCACTGCACTTTCACCCTGG 2382

Qy    2297 AGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGG 2356
           || |   |||| |  | | ||| | |||| ||| ||| | || |||||| |||||| || |
Db    2383 AGAGACATAGCCTAGCCTCCACTGAGTTCACCTGTAAGGTCTGCGTGCGGCAGGTAGAAG 2442

Qy    2357 GCGACGGGCAGAGCTTCAGCATCAACTTCAACATCAC---CAAGGACACAAGGTTTGCTG 2413
           | || || |||| |||  |||     |  || ||  |  |  |    | |  | |   |
Db    2443 GGGAAGGCCAGATTTTCCAGCTGCACACCACGCTGGCTGAGACGCCTGCTGGCTCCCTGG 2502

Qy    2414 AGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTCA 2473
           |   ||      |    ||     | | | | | | |||| ||| ||||||||
Db    2503 ATGCACTCTGCTCTGCCCCTGGCAATGCTGCCACCACACAGCTGGGACCCTATGCCTTCA 2562

Qy    2474 AGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGCGGG 2533
           |||| ||  |   ||| || |||||||| ||||||||| | ||| |||    ||||
Db    2563 AGATACCACTGTCCATCCGCCAGAAGATCTGCAACAGCCTGGACGCCCCCAACTCACGGG 2622

Qy    2534 GTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCTTTG 2593
           |   ||||||||||    |||| ||||| |||  | |||||| | |  | || | || |
Db    2623 GCAATGACTGGCGGCTGTTGGCACAGAAGCTCTCCATGGACCGGTACCTGAACTACTTCG 2682

Qy    2594 CCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCCCCA 2653
           || ||||  | || ||||||| | ||||| | ||||| ||||| | |   ||
Db    2683 CCACCAAAGCTAGTCCCACAGGCGTGATCTTAGACCTCTGGGAAGCTCGGCAGCAGGATG 2742

Qy    2654 ACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTGGCC 2713
           | || |||||| |  ||||| ||  |||  |  |||||| ||    ||
Db    2743 ATGGGGACCTCAACAGCCTGGCCAGTGCCTTGGAGGAGATGGGCAAGAGTGAGATGCTGG 2802

Qy    2714 TCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
           |   ||     | || | || |||||||
Db    2803 TAGCCATGACCACTGATGGCGATTGCTGA 2831
```

RESULT 7
US-09-949-016-4794
; Sequence 4794, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

```
  Query Match                30.3%;   Score 833.6;   DB 4;   Length 3008;
  Best Local Similarity      62.1%;   Pred. No. 7.4e-177;
  Matches 1435;  Conservative      0;  Mismatches  814;   Indels    63;   Gaps      5;


Qy        485 TGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCG 544
              I II IIII  II IIIIIIII II II I  IIIII IIIII  IIII IIII   II
Db         10 TACGGAAGACATTTGAGCAGGAACCCCTAGGAAAGGAAGTGTCTTTGGAACAGGAAGTCT 69


Qy        545 TGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACG 604
              I II I  II II IIIII II II IIIII    II IIIIIIIII III I    II I
Db         70 TACTCCAGTGTCGACCACCTGAAGGGATCCCAGTGGCTGAGGTGGAATGGTTGAAAAATG 129


Qy        605 AGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGG 664
              I IIII I I II II         II I  III I II II II   II IIII III
Db        130 AAGACATAATTGATCCCGTTGAAGATCGGAATTTTTATATTACTATTGATCACAACCTCA 189


Qy        665 TGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCG 724
              I  I     IIIIIIIII II  IIII II II II IIIIIIII II IIIII IIIII I
Db        190 TCATAAAGCAGGCCCGACTCTCTGATACTGCAAATTACACCTGTGTTGCCAAAAACATTG 249


Qy        725 TGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGA 784
              I II     I    II I I II IIIIIIII IIIII II IIIIIIIII IIIII I
Db        250 TTGCCAAGAGGAAAAGTACAACTGCCACTGTCATAGTCTATGTCAACGGTGGCTGGTCCA 309


Qy        785 CGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCC 844
              I IIIIII IIIIIIII II II I I  I IIIIIIII II II I  IIIIIIII I
Db        310 CCTGGACGGAGTGGTCTGTGTGTAACAGCCGCTGTGGACGAGGGTATCAGAAACGTACAA 369


Qy        845 GGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCC 904
              III  II IIIIIIIIIII II IIIII IIIII II IIIIIIIII IIIIIII III I
Db        370 GGACTTGTACCAACCCGGCACCACTCAATGGGGGTGCCTTCTGTGAAGGGCAGAGTGTGC 429


Qy        905 AGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGT 964
              IIIIII IIIIII  I II  I IIIIIIIII II IIIII IIII  II IIIIIIIIII
Db        430 AGAAAATAGCCTGTACTACGTTATGCCCAGTGGATGGCAGGTGGACGCCATGGAGCAAGT 489


Qy        965 GGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCAC 1024
              IIII I IIIII  II IIIIIIIIIIIIIIII II I IIIIII I I  IIIII I
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```
Db     490  GGTCTACTTGTGGAACTGAGTGCACCCACTGGCGCAGGAGGGAGTGCACGGCGCCAGCCC  549

Qy    1025  CCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTG 1084
             ||    || |||||  |||| |||  |  |||   | | ||  | ||   ||||| ||   ||
Db     550  CCAAGAATGGAGGCAAGGACTGCGACGGCCTCGTCTTGCAATCCAAGAACTGCACTGATG  609

Qy    1085  ACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGG---CCTCA 1141
             || ||  |  | || | |||| |||  | || ||  || || ||||||||| ||     | |
Db     610  GGCTTTGCATGCAGACTGCTCCTGATTCAGATGATGTTGCTCTCTATGTTGGGATTGTGA  669

Qy    1142  TCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGA 1201
             | || |||  ||| |||||||    |    | |||| || | | || ||| |    || 
Db     670  TAGCAGTGATCGTTTGCCTGGCGATCTCTGTAGTTGTGGCCTTGTTTGTGTATCGGAAGA  729

Qy    1202  AGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTCCAGC 1261
             | |    |   | || |||||| |   |||||| ||    |||| ||     ||||| ||||
Db     730  ATCATCGTGACTTTGAGTCAGATATTATTGACTCTTCGGCACTCAATGGGGGCTTTCAGC  789

Qy    1262  CCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCGGACC 1321
             | || | |||||||        |||| |   | ||||||| |  | | · || ||||
Db     790  CTGTGAACATCAAG---------GCAGCAAGACAAGATCTGCTGGCTGTACCCCCAGACC  840

Qy    1322  TCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATG-----GGCC 1376
             |||   |  |  |  | ||| |||   ||  | ||| | ||| | ||| || ||       | |
Db     841  TCACGTCAGCTGCAGCCATGTACAGAGGACCTGTCTATGCCCTGCATGACGTCTCAGACA  900

Qy    1377  CAGCCCCAAGTTCCAGCTCACCAAT----GGGCACCTGCTCAGCCCCCTGGGTGGCGGCC 1432
             | ||||| ||| ||| ||| |||||      | | | |||| || ||          |
Db     901  AAATCCCAATGACCAACTCTCCAATTCTGGATCCACTGCCCAACCTGAAAATCAAAGTGT  960

Qy    1433  GCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCT 1492
             | ||||      | | ||| ||| |  | ||||| |||    ||| ||  |||
Db     961  ACAACACCTCAGGTGCTGTCACCCCCCAAGATGACCTCTCTGAGTTTACGTCCAAGCTGT 1020

Qy    1493  CCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGA------------ 1540
             || | |||| |       |   | | ||    |     | || || | |||
Db    1021  CCCCTCAGATGACCCAGTCGTTGTTGGAGAATGAAGCCCTCAGCCTGAAGAACCAGAGTC 1080

Qy    1541  -----------------------------CCTATGGGACCTTCAACTTCCTCGGGGGCC 1570
                                          | | ||| | ||||||||| || || || |
Db    1081  TAGCAAGGCAGACTGATCCATCCTGTACCGCATTTGGCAGCTTCAACTCGCTGGGAGGTC 1140

Qy    1571  GGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAG 1630
             || || | || ||| ||||  ||||| | || || || ||| | |  ||||| |||| ||
Db    1141  ACCTTATTGTTCCCAATTCAGGAGTCAGCTTGCTGATTCCCGCTGGGGCCATTCCCCAAG 1200

Qy    1631  GGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAG 1690
             |||    |||| ||  ||  | |||  |  | |||| |       |||     |||||  |
Db    1201  GGAGAGTCTACGAAATGTATGTGACTGTACACAGGAAAGAAACTATGAGGCCACCCATGG 1260

Qy    1691  CTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCA 1750
             || || |||||||| || |||  |||   || |  | || ||||||||| |||||| | |  |||||||
Db    1261  ATGACTCTCAGACACTTTTGACCCCTGTGGTGAGCTGTGGGCCCCCAGGAGCTCTGCTCA 1320

Qy    1751  CCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCC 1810
             |||| ||||||| |||| ||||| | |||||| | ||||| |  || |||  |  ||||| |
Db    1321  CCCGCCCAGTCGTCCTCACTATGCATCACTGCGCAGACCCCAATACCGAGGACTGGAAAA 1380
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Qy   1811 TGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGG 1870
          | |  ||||| || ||| |     ||||   ||||||||||||| ||    | || ||||
Db   1381 TACTGCTCAAGAACCAGGCAGCACAGGGACAGTGGGAGGATGTGGTGGTGGTCGGGGAGG 1440

Qy   1871 AGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCG 1930
          |       | || || || |||| |||||||| ||     |||||| || || |||| |
Db   1441 AAAACTTCACCACCCCCTGCTACATTCAGCTGGATGCAGAGGCCTGCCACATCCTCACAG 1500

Qy   1931 AGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCC 1990
          || | || || || |||||||| ||| | || || | || || || || |||||||
Db   1501 AGAACCTCAGCACCTACGCCCTGGTAGGACATTCCACCACCAAAGCGGCTGCGAAGCGCC 1560

Qy   1991 TCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCT 2050
          |||||||     | |||| ||| ||   |||| |||| || |||||||| ||||||  ||||
Db   1561 TCAAGCTGGCCATCTTTGGGCCCCTGTGCTGCTCCTCGCTGGAGTACAGCATCCGAGTCT 1620

Qy   2051 ACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGG 2110
          |||| ||| ||||||||| |||||| || |||||     |  | || || |||| ||| |||
Db   1621 ACTGTCTGGATGACACCCAGGATGCCCTGAAGGAAATTTTACATCTTGAGAGACAGATGG 1680

Qy   2111 GGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGC 2170
          | ||||||||   |  | || ||   || || || || || | |||    |||||||||||
Db   1681 GAGGACAGCTCCTAGAAGAACCTAAGGCTCTTCATTTTAAAGGCAGCACCCACAACCTGC 1740

Qy   2171 GCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACC 2230
          |||| || ||  ||||||  |  |||   | ||||| |||||||| || | ||| | || |
Db   1741 GCCTGTCAATTCACGATATCGCCCATTCCCTCTGGAAGAGCAAATTGCTGGCTAAATATC 1800

Qy   2231 AGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCA 2290
          |||| || || ||||| ||  | |||| |||  | || |  |  || |||||||||||||||
Db   1801 AGGAAATTCCATTTTACCATGTTTGGAGTGGATCTCAAAGAAACCTGCACTGCACCTTCA 1860

Qy   2291 CCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGG 2350
          | ||||| |   | | |||| | ||| |     || |||| ||||| || || ||| ||||||
Db   1861 CTCTGGAAAGATTTAGCCTGAACACAGTGGAGCTGGTTTGCAAACTCTGTGTGCGGCAGG 1920

Qy   2351 TGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGACACAAGGTTTG 2410
          |||| || || || |||||||| |||    |||||| || | | | |||| | ||
Db   1921 TGGAAGGAGAAGGGCAGATCTTCCAGCTCAACTGCACCGTGTCAGAGGAACCTACTGGCA 1980

Qy   2411 CTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCT 2470
          ||  ||| |     ||||| ||    || | | |    ||| |||||||||||| |
Db   1981 TCGATTTGCCGCTGCTGGATCCTGCGAACACCATCACCACGGTCACGGGGCCCAGTGCTT 2040

Qy   2471 TCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGC 2530
          |||  ||||| |||  || || ||||||||||||| |    |||||||||| | |||   | |
Db   2041 TCAGCATCCCTCTCCCTATCCGGCAGAAGCTCTGTAGCAGCCTGGATGCCCCCAGACGA 2100

Qy   2531 GGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCT 2590
          | ||     |||||| ||| ||| |||||||||| || || |||||||||| | | | | ||
Db   2101 GAGGCCATGACTGGAGGATGCTGGCCCATAAGCTGAACCTGGACAGGTACTTGAATTACT 2160

Qy   2591 TTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCC 2650
          ||||| ||||     ||||||| || | | | | ||||| | || ||||| || | | ||||||
Db   2161 TTGCCACCAAATCCAGCCCAACTGGCGTAATCCTGGATCTTTGGGAAGCACAGAACTTCC 2220
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Qy    2651 CCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTG 2710
           |  | || ||||||  |||   |||||| || |   ||| | | ||||    |  || | |
Db    2221 CAGATGGAAACCTGAGCATGCTGGCAGCTGTCTTGGAAGAAATGGGAAGACATGAAACGG 2280

Qy    2711 GCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
           | | |  ||   | ||| |   ||| | |
Db    2281 TGGTGTCCTTAGCAGCAGAAGGGCAGTATTAA 2312


RESULT 8
US-09-969-532-9
; Sequence 9, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;   APPLICANT: Walke, D. Wade
;   APPLICANT:  Scoville, John
;   TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
;   FILE REFERENCE: LEX-0244-USA
;   CURRENT APPLICATION NUMBER: US/09/969,532
;   CURRENT FILING DATE:  2001-10-02
;   PRIOR APPLICATION NUMBER: US 60/237,280
;   PRIOR FILING DATE: 2000-10-02
;   NUMBER OF SEQ ID NOS: 33
;   SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 9
;   LENGTH: 2736
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-9

   Query Match              17.7%;   Score 487;   DB 4;   Length 2736;
   Best Local Similarity    50.9%;   Pred. No. 2.7e-99;
   Matches 1305;  Conservative     0;  Mismatches 1230;  Indels    30;  Gaps     5;

Qy     172 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 231
           || || || |||  |  | ||||||| || ||||  || || ||||||| ||| || |
Db     157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216

Qy     232 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 291
           ||    ||||||| ||   ||| |||| |||||| |||||||| ||||||||| ||||||||
Db     217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276

Qy     292 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 351
           |  |||  || |||||     ||   | |     |||| ||||  || ||      |
Db     277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC .336

Qy     352 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 411
           || ||    ||| |||||| | ||||| ||||| ||| |  |    |||| ||||| ||
Db     337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396

Qy     412 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 471
           |||||||||||| ||||| ||||| |   ||| ||| ||||||||    |||||||| |
Db     397 TGGTGCCAGTGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456

Qy     472 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 531
           ||||||||| || || || |||||| || || || || ||| | | | ||| || || |
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Db    457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516

Qy    532 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 591
              ||       || || |||| |||||| |||||| |||||   |||||| | |||||||||||
Db    517 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576

Qy    592 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 651
              |||||      || || || |    | |||| |      |  |||     || |   ||| || |
Db    577 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636

Qy    652 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 711
              || || | |||| |   |   | |||||| || ||   | ||| | |   || |||||||||| ||
Db    637 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696

Qy    712 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 771
              ||       ||||||||||||      |  | |||    || ||    |||| |    ||||||||||| |
Db    697 GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 756

Qy    772 GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 831
              ||   | |||         ||||| ||| |||||||||||||||| |       ||||
Db    757 GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTGA--------A 807

Qy    832 CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 891
              ||     |||| ||||    ||||| |     || | ||    || ||||||   ||||||||
Db    808 CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 867

Qy    892 GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 951
              ||| |    |        |  | |||| |                  |     | |
Db    868 GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGATAAAAAACCT 927

Qy    952 CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 1011
              |         |        |           |  |  || | ||                    | |||
Db    928 CTTCATGAAATAAAACCCCAAAGCATTGAGAATGCCAGCGACATTGCTTTGTACTCGGGC 987

Qy   1012 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1071
              | |   | || |        || | |     | |    | |    || |||   ||| ||| ||
Db    988 TTGGGTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACCCTTTACAGACGG 1047

Qy   1072 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1131
              | |    |        |   |   |   || || ||| |    |   |  | | |  || |
Db   1048 AGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACAGGTGGCTTCCA- 1106

Qy   1132 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1191
              | | ||| | |      | ||| | ||| || |   | ||| ||   || || |
Db   1107 --GACCTTCAACTTCAAAACAGTCCGTCAAGCCAAGAATATCATGGAACTAATGATACAA 1164

Qy   1192 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1251
              |             |   ||| |   ||  ||    | |   ||| ||
Db   1165 GAAAAATCCTTTGGTAACTCCCTGCTCCTGAATTCTGCCATGCAGCCAGATCTGACAGTG 1224

Qy   1252 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1311
              ||          | | | ||          | |              | |   |||||| |
Db   1225 AGCCGGACATACAGCGGACCCATCTGTCTGCAGGACCCTCTGGACAAGGAGCTCATGACA 1284

Qy   1312 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1371
              ||| |     |||  | | |      | | || | || | ||       | |
Db   1285 GAGTCCTCACTCTTTAACCCTTTGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATG 1344
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Qy    1372 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1431
           |  ||        ||    ||   |  |    |||      |   |          |
Db    1345 GTTTCCCTGGGAGTGTCTGAGAGAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT 1404

Qy    1432 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1491
           | |||   |    |||||||||| |  ||      |  ||     |    |  || | || 
Db    1405 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1464

Qy    1492 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1551
            |  |  |            || || |||  | | ||      | |||   |||     |
Db    1465 CAAAATCTGTC---------ATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1515

Qy    1552 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1611
           ||   |    | ||||| ||  |  | || || ||||||||| | |||  | ||||| ||
Db    1516 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1575

Qy    1612 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1671
           | | |||||||| ||    | |||   | |||||| || | | | | |   |||  |  |
Db    1576 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1632

Qy    1672 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1731
           ||     ||  | |    || |||||| | ||   || || |||||||||     || | ||||||
Db    1633 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1692

Qy    1732 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1791
           || || | |  |    | |||||   ||  |    || | ||    |||||||  ||   |
Db    1693 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1752

Qy    1792 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1851
           ||  ||||    ||||    | |  |  | || ||    ||      |||||| ||||||||
Db    1753 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1812

Qy    1852 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1911
           ||| ||   ||| || ||  |    | ||     ||  |||||||    |||| ||   |
Db    1813 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1866

Qy    1912 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1971
           || ||  | ||  ||   ||     | ||   | || || ||| |      ||||| |  |||
Db    1867 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1926

Qy    1972 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 2031
           |     |||||  ||| |||| ||  || || |    | ||||      || ||||| | ||||||
Db    1927 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1986

Qy    2032 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2091
           || ||||||| |   | || ||||| ||| | | ||||     |||| | |||| |||||||
Db    1987 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2046

Qy    2092 CAGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2151
           || | ||| |  || ||||||||| |  | || |||      | |||| |||||  |||||
Db    2047 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2106

Qy    2152 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2211
           | |   |  | |    |    || |   | ||  ||| ||| ||   | ||| |||  | |
Db    2107 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2166
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Qy   2212 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2271
          || |    |    || |||||| |||| || |  | | | ||| | | | ||| |
Db   2167 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2226

Qy   2272 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2331
          | ||||||| ||||| |||||||||||  |  |||| || |  | ||| |||||
Db   2227 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2286

Qy   2332 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2391
          || | || |  ||||| |  | ||| || |||| | ||   | |          |||
Db   2287 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2346

Qy   2392 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2451
          ||     | |    |       |        |         | ||     |||| ||
Db   2347 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2406

Qy   2452 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2511
          | |    ||||||| ||||||||| || |||| |  ||| | ||| ||| |  |
Db   2407 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2466

Qy   2512 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2571
          | || | ||| |    ||        ||||||| ||  | || |||||| |  | |
Db   2467 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2526

Qy   2572 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2631
          |||| || |      | || || | |    ||||| | || | || | ||||||
Db   2527 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTGAACCTG 2586

Qy   2632 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2691
          ||||| || || ||    | | || | || |  | ||||| || | |
Db   2587 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2646

Qy   2692 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2736
          | ||   | || || |   |||| |  | || || |  ||
Db   2647 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2691
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RESULT 9
US-09-969-532-33
; Sequence 33, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
; APPLICANT: Walke, D. Wade
; APPLICANT: Scoville, John
; TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
; FILE REFERENCE: LEX-0244-USA
; CURRENT APPLICATION NUMBER: US/09/969,532
; CURRENT FILING DATE: 2001-10-02
; PRIOR APPLICATION NUMBER: US 60/237,280
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 33
;   LENGTH: 3411
;   TYPE: DNA
;   ORGANISM: homo sapiens

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Query Match              17.7%;  Score 487;  DB 4;  Length 3411;
Best Local Similarity    50.9%;  Pred. No. 2.9e-99;
Matches 1305;  Conservative    0;  Mismatches 1230;  Indels   30;  Gaps    5;


Qy      172 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 231
            || || ||  ||| | | ||||||  || ||||  || ||  ||||||  ||| ||   |
Db      271 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 330


Qy      232 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 291
            ||   ||||||  ||   ||| ||||| |||||||| |||||||| ||||||||| |||||
Db      331 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 390


Qy      292 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 351
            |  |||   ||  |||||   ||   |    |||| |||| ||  || |           |
Db      391 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 450


Qy      352 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 411
            || ||   ||| |||||   | ||||| ||||| |||  |      |||| ||||| ||
Db      451 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 510


Qy      412 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 471
            ||||||||||| ||||| ||||||  |   ||| ||| |||||||    ||||||||   |
Db      511 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 570


Qy      472 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 531
            |||||||||  || || || || ||||| || ||  || || || |  | | ||| ||  |
Db      571 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 630


Qy      532 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 591
            ||         || || |||| |||||| |||||  ||||| ||||| | ||||||||||||
Db      631 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 690


Qy      592 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 651
            |||||     || || || || | |   | ||| |    | |||   || | ||| || |
Db      691 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 750


Qy      652 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 711
            || || | ||||| |  | | | ||||| || ||  | ||| | | || |||||||||| ||
Db      751 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 810


Qy      712 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 771
            ||       |||||||||  | | ||   ||   || || |||| | ||||||||||| |
Db      811 GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 870


Qy      772 GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 831
            ||  |  ||     |||| ||| ||| |||||||||||||  |     ||||
Db      871 GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTGA---------A 921


Qy      832 CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 891
            ||     |||| |||     |||||  || || | ||   || ||||||    ||||||||
Db      922 CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 981


Qy      892 GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 951
            ||| |   |    |   |   |||| |     |||| |         |   |   |
Db      982 GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGATAAAAAACCT 1041
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Qy      952 CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 1011
             |     |      |      |      | |   || | ||                | ||
Db     1042 CTTCATGAAATAAAACCCCAAAGCATTGAGAATGCCAGCGACATTGCTTTGTACTCGGGC 1101

Qy     1012 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1071
             | |   | ||   |     || | |    | |  ||    ||| ||| ||
Db     1102 TTGGGTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACCCTTTACAGACGG 1161

Qy     1072 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1131
             | |   |     |   |   |     || || |||   |   |   | | ||| |
Db     1162 AGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACAGGTGGCTTCCA- 1220

Qy     1132 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1191
             | | ||| |  |   | ||| | | | | |   |   || || |
Db     1221 --GACCTTCAACTTCAAAACAGTCCGTCAAGCCAAGAATATCATGGAACTAATGATACAA 1278

Qy     1192 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1251
             |           | |   ||| | || || | | ||  || |  || ||
Db     1279 GAAAAATCCTTTGGTAACTCCCTGCTCCTGAATTCTGCCATGCAGCCAGATCTGACAGTG 1338

Qy     1252 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1311
             ||         | | |  || | |   || |      | | | |||||  |
Db     1339 AGCCGGACATACAGCGGACCCATCTGTCTGCAGGACCCTCTGGACAAGGAGCTCATGACA 1398

Qy     1312 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1371
             || |   ||| | | | |     | | | | | | | ||    | |
Db     1399 GAGTCCTCACTCTTTAACCCTTTGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATG 1458

Qy     1372 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1431
             |   ||        ||   ||      | |    |||     | |         |
Db     1459 GTTTCCCTGGGAGTGTCTGAGAGAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT 1518

Qy     1432 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1491
             | |||   |   |||||||||| |   ||     | ||     |      | || | ||
Db     1519 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1578

Qy     1492 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1551
             |   |         || || |||  | | ||        | |||    ||| |
Db     1579 CAAAATCTGTC---------ATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1629

Qy     1552 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1611
             ||   |     | |||||  ||   |   |  || || |||||||||  | ||| | ||||| ||
Db     1630 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1689

Qy     1612 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1671
             | | ||||||| ||     | |||     | |||||| ||  | | |   ||| |  |
Db     1690 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1746

Qy     1672 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1731
             ||     ||   |  |     || |||||| | ||   || |||||||||    || | ||||||
Db     1747 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1806

Qy     1732 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1791
             || || | | |   |  ||||| ||  |   || | ||  |||||||  || |
Db     1807 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1866
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Qy   1792 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1851
          ||  ||||    ||||   |  |  | || |||  | |      ||||||  ||||||||
Db   1867 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1926

Qy   1852 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1911
          ||| ||   |||  || ||  |  | ||        ||  |||||||| ||||  || |
Db   1927 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1980

Qy   1912 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1971
          || ||  | ||  ||   ||    | ||  | || || ||| ||  ||||||| |  |||
Db   1981 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 2040

Qy   1972 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 2031
          |    |||||| |||| ||  || |||  |    ||||     || |||| | ||||||
Db   2041 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 2100

Qy   2032 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2091
          || |||||| |  | || ||||| || || |   ||||    |||| | ||||| |||||
Db   2101 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2160

Qy   2092 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2151
          || | ||| |   || || |||||||||| |  | ||| ||     |||||| ||||||
Db   2161 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2220

Qy   2152 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2211
          |  |  |  | | | |  ||  | ||    |||   ||| |  ||   | ||| ||  |
Db   2221 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2280

Qy   2212 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2271
          || |    |   || |||||| |||| |||| | | | | ||    | || | |  | |
Db   2281 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2340

Qy   2272 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2331
           | |||||||  |||||| ||||||||| ||   | ||||  | |   |   | ||||||
Db   2341 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2400

Qy   2332 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2391
          ||  | ||  |   ||||| |  | ||| | |  ||||  | | |   |   |    |||
Db   2401 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2460

Qy   2392 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2451
           ||   || |  | |    |  | |   | |  | |     ||  | ||||   |||| ||
Db   2461 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2520

Qy   2452 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2511
          | |   |||||||| ||||||||| |||||||| || |||| |  | |||  |||  | |
Db   2521 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2580

Qy   2512 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2571
          | || | | | |||  |   ||    |||||||| || |||||||| |  | | | |
Db   2581 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2640

Qy   2572 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2631
          ||||   || | |  | |  | |  | | |     |||||  |  | | | | ||||||||
Db   2641 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2700

Qy   2632 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2691
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             |||||  || || ||      |   | ||  | ||     |   |||||    ||  | |  |
Db    2701 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2760   .


Qy    2692 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2736
             | ||    | || || |     |||| |   | || || |  ||
Db    2761 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2805
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RESULT 10
US-09-969-532-11
; Sequence 11, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
; APPLICANT: Walke, D. Wade
; APPLICANT: Scoville, John
; TITLE OF INVENTION: No. 6777232e1 Human Membrane Proteins and Polynucleotides
Encoding the Same
; FILE REFERENCE: LEX-0244-USA
; CURRENT APPLICATION NUMBER: US/09/969,532
; CURRENT FILING DATE: 2001-10-02
; PRIOR APPLICATION NUMBER: US 60/237,280
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 11
;   LENGTH: 2703
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-11

  Query Match   17.0%;   Score 467;   DB 4;   Length 2703;
  Best Local Similarity   51.0%;   Pred. No. 8.1e-95;
  Matches 1307; Conservative   0; Mismatches 1195; Indels   63; Gaps   6;

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Qy     172 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 231
             || || ||  ||| | | ||||||| || ||||  || ||  ||||||| ||| || |
Db     157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216


Qy     232 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 291
             ||    ||||||| ||     ||| ||||  |||||| |||||||||| ||||||||| ||||||||
Db     217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276


Qy     292 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 351
             |  |||   || |||||      ||   |     |||| ||||  || ||          |
Db     277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 336


Qy     352 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 411
             || ||     ||| |||||  | |||||| ||||| ||| |  |     |||| ||||| ||
Db     337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396


Qy     412 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 471
             |||||||||| ||||| |||||| |   ||| ||| |||||||    |||||||| |
Db     397 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456


Qy     472 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 531
             ||||||||| ||  ||  || || |  ||  ||  |   || || || |  |   | ||| |
Db     457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516
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Qy    532  GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 591
               ||      || || |||| |||||| |||||| |||||   |||||  | ||||||||||||
Db    517  GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576

Qy    592  TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 651
               |||||    || || || |  | |||| |    |  ||| |  || |   ||| || |
Db    577  TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636

Qy    652  GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 711
               || || | |||| |  | | |||||| || ||  | ||| | |  || |||||||||| ||
Db    637  GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696

Qy    712  GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 771
               ||      ||||||||||      |  | |||    || ||  |||| |  ||||||||| |
Db    697  GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 756

Qy    772  GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 831
               || | |||     |||||  |||  |||||||||||||||||||||  |     ||||
Db    757  GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTG---------AA 807

Qy    832  CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 891
               ||     |||| ||||    |||||     || | ||   || |||||||   ||||||||
Db    808  CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 867

Qy    892  GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 951
               || | |        |     | ||||        |  ||| |   |||| || ||
Db    868  GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGGCATTGAGAAT 927

Qy    952  CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 1011
               |  |   ||    | |  || || ||   |||            ||||
Db    928  GCCAGCGACATTGCTTTGTACTCGGGCTTGG----------------------GTGC 962

Qy   1012  TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1071
               |     | | | | |||     || |   ||| ||     |   |||
Db    963  TGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACCCTTTACAGACGGA------- 1015

Qy   1072  AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1131
                  |   |||||| |  | ||  |  | |        |||          |       ||
Db   1016  ----GCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACAGGTGGCTTC 1071

Qy   1132  GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1191
               | | ||| |  |   | ||| | | |  | |   |   |  | || ||  |
Db   1072  CAGACCTTCAACTTCAAAACAGTCCGTCAAGCCAAGAATATCATGGAACTAATGATACAA 1131

Qy   1192  TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1251
               |          |   ||| |    ||   ||  ||  | |    |||  |
Db   1132  GAAAAATCCTTTGGTAACTCCCTGCTCCTGAATTCTGCCATGCAGCCAGATCTGACAGTG 1191

Qy   1252  GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1311
               ||         |  |  |  ||         | |       | |  |||||  |
Db   1192  AGCCGGACATACAGCGGACCCATCTGTCTGCAGGACCCTCTGGACAAGGAGCTCATGACA 1251

Qy   1312  CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1371
               ||  |   |||   |  | |     | |   |     | || |    |  |
Db   1252  GAGTCCTCACTCTTTAACCCTTTGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATG 1311
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Qy   1372 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1431
          |  ||          ||  ||   |  |    |||      |  |           |
Db   1312 GTTTCCCTGGGAGTGTCTGAGAGAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT 1371

Qy   1432 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1491
          | |||    |    |||||||||| |  ||     |  ||   |    |  | |  | ||
Db   1372 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1431

Qy   1492 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1551
           |  |          ||  || |||  |  |  ||        |  |||   |||   |
Db   1432 CAAAATCTGTC---------ATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1482

Qy   1552 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1611
          ||   |     |  ||||| ||  |   |  || |||||||||  | |||  | |||||  ||
Db   1483 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1542

Qy   1612 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1671
          |  | ·||||||  ||    | |||     | |||||  ||   |   |  ||| |  |
Db   1543 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1599

Qy   1672 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1731
          ||    ||  | |   ||  |||| | ||  ||  |||||||||    ||  |  |||||||
Db   1600 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1659

Qy   1732 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1791
          || || | |   | |||||    || |   |  |   | |||   |||||||  || |
Db   1660 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1719

Qy   1792 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1851
          ||  ||||   ||||    || | |   | || |||   |      ||||||  |||||||||
Db   1720 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1779

Qy   1852 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1911
          ||| ||   ||| || || |  |  | ||     ||  |||||||  |||| || |
Db   1780 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1833

Qy   1912 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1971
          || ||  | ||  ||   ||   ||  | || || ||  | |||||| |   |||
Db   1834 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1893

Qy   1972 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 2031
          |     ||||| |||| || ||| || |    | |||      || |||| | ||||||
Db   1894 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1953

Qy   2032 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2091
          || |||||| |  | || ||||| || | | |||| ||| | |||  | |||| |||||
Db   1954 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2013

Qy   2092 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2151
          || | ||| | || |||||||| | | | || |||   |  ||||| |||||
Db   2014 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2073

Qy   2152 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2211
          | | | |   |  ||   |  | || |  || |||   | ||  | ||| |||| ||  |
Db   2074 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2133

Qy   2212 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2271
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                 || |      |        || ||||||  |||| || |  | | | |||    | |  | || |
Db    2134 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2193

Qy    2272 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2331
                 | |||||||  ||||| |||||||||||  |  |||| || |   | ||| |||||
Db    2194 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2253

Qy    2332 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2391
                 || | || |   ||||| | | ||| | |  |||| | ||    | |       ||| 
Db    2254 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2313

Qy    2392 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2451
                  ||     | |      |        |         |    ||       |||| ||
Db    2314 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2373

Qy    2452 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2511
                 | |    ||||||| |||||||| || |||| .|  ||| | ||| |||   | | |
Db    2374 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2433

Qy    2512 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2571
                 | || | ||| |        ||      ||||||| ||  | || ||||||  | | |
Db    2434 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2493

Qy    2572 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2631
                 ||||  || |       | || ||  | |     ||||| | || | ||  | ||||||
Db    2494 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2553

Qy    2632 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2691
                 ||||| || || ||     |  | ||  | ||    |  |||||     || | | |
Db    2554 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2613

Qy    2692 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2736
                 | ||    | || || |    |||| |   | || ||  | || 
Db    2614 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2658


RESULT 11
US-09-969-532-13
; Sequence 13, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;  APPLICANT: Walke, D. Wade
;  APPLICANT:  Scoville, John
;  TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
;  FILE REFERENCE: LEX-0244-USA
;  CURRENT APPLICATION NUMBER: US/09/969,532
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US 60/237,280
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 33
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13
;   LENGTH: 2694
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-13
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  Query Match              16.7%;  Score 460.2;  DB 4;  Length 2694;
  Best Local Similarity    51.9%;  Pred. No. 2.7e-93;
  Matches 1330;  Conservative    0;  Mismatches 1163;  Indels  72;  Gaps    10;

Qy      172 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 231
                || || || ||| |  | ||||||| || |||| || ||  ||||||  ||| ||  |
Db      157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216

Qy      232 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 291
                ||    ||||||  ||      ||| |||| |||||| ||||||||| ||||||||| ||||||||
Db      217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276

Qy      292 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 351
                |   |||    ||  ||||||       ||      |      |||| ||||   ||  ||         |
Db      277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 336

Qy      352 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 411
                || ||   ||| |||||| | |||||| ||||| ||| | |      |||| ||||| ||
Db      337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396

Qy      412 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 471
                |||||||||||| |||||| |||||| |  | ||| ||| |||||||   ||||||||    |
Db      397 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456

Qy      472 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 531
                |||||||||   || || || || ||||| || || || || || |  |  | ||| ||  || |
Db      457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516

Qy      532 GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 591
                ||          || || |||| |||||| ||||| ||||| |||||  ||||| | ||||||||||||
Db      517 GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576

Qy      592 TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 651
                |||||     || || || || |   | ||| |   | |||  || | ||| || |
Db      577 TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636

Qy      652 GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 711
                || || | |||| |  |  | |||||| || ||  | ||| |  | || |||||||||| ||
Db      637 GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696

Qy      712 GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 771
                ||       ||||||||||     |  | |||  || || |||| |  | |||||||||| |
Db      697 GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 756

Qy      772 GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 831.
                || | |||     ||||| ||| ||||||||||||||||| |     ||||
Db      757 GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTG---------AA 807

Qy      832 CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 891
                ||       |||| ||||     |||||       || | ||   || ||||||   ||||||||
Db      808 CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 867

Qy      892 GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 951
                ||| |    | | |   ||| ||      || | || |   | | | |      | |
Db      868 GGTC--TAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGATAAAAAAC 925
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Qy     952 CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 1011
            |  |  |  || |     || |  ||     |   | |  ·  |    |  |  |+ |
Db     926 CTCTTCATGAAATAAAACCCCAAAGCATTGAG----AATGCCAGCGACATTGCTTTGTAC 981

Qy    1012 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1071
            || | |    |   | |    || |  |  |    || || || ||  |       |  |
Db     982 TCGGGCTTGGGTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGT------CACC 1035

Qy    1072 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1131
             |  |  |   || |   |||   |  |     | |   ||  |||      || |
Db    1036 CTTTACAGACGGAGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACA 1095

Qy    1132 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1191
            |  ||| ||     || |  | ||      |||| |        ||| ||||       |
Db    1096 GGTGGCTTCCAGACCTTCAACTTCAAAACAGTCCGTCAAGGTAACTCCCTGCTCCTGAAT 1155

Qy    1192 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1251
            |  |  |        |||  | |||| ||| |    || |      |     | | ||
Db    1156 TCTGCCATGCA--------GCCAGATCTGACAGTGAGCCGGACATACAGCGGACCCATCT 1207

Qy    1252 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1311
            | || |  | || || | |  |  || ||  | |   ||  | | |  ||  ||    |
Db    1208 GTCTGCAGGACCCTCTGGACAAGG---AGCTCATGACAGAGTCCTCACTCTTTAACCCTT 1264

Qy    1312 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1371
             |  |       |     |    | |  ||| |    |  || |||    | | |
Db    1265 TGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATGGTTTCCCTGGGAGTGTCTGAGA 1324

Qy    1372 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1431
            | || || ||| |       ||| |    || |   |
Db    1325 GAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT--------------------- 1362

Qy    1432 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1491
            | ||| |     | |||||||||| | | ||  |  || | | | || | | || | ||
Db    1363 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1422

Qy    1492 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1551
             |  |  |         || || ||| | | || |   | ||| ||| ||| |
Db    1423 CAAAATCTG--------TCATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1473

Qy    1552 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1611
            ||   |   | ||||| || | |  |  || || |||||||||| | ||| | |||||||
Db    1474 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1533

Qy    1612 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1671
            | | ||||||| ||    |··|·|| |     ||||| || |||  | |   ||| |  |
Db    1534 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1590

Qy    1672 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1731
            ||    ||  ||  |    || ||||| | || || |||||||||     || | ||||||
Db    1591 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1650

Qy    1732 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1791
            || || | || | ||  ||||| |   || |   || | ||||||||||  ||  |
Db    1651 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1710

Qy    1792 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1851
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                          ||  ||||     ||||   | |   | || |||  | |     ||||||  |||||||||
Db       1711 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1770

Qy       1852 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1911
                          ||| ||    ||| || || |  | ||     ||  ||||||||   |||| || |
Db       1771 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1824

Qy       1912 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1971
                          || ||  | ||  ||    ||    | ||    || ||| ||   |||||| |  |||
Db       1825 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1884

Qy       1972 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 2031
                          |     ||||| ||||   || ||| ||    | ||||     || |||| | ||||||
Db       1885 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1944

Qy       2032 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2091
                          || |||||| | | || ||||| || |  | |||| |||| |    |||| |||||
Db       1945 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT 2004

Qy       2092 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2151
                          || | ||| |   || |||||||||  |  | || |||      | |||||| |||||
Db       2005 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2064

Qy       2152 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2211
                          |  | | |   | | ||| |  | ||  ||| |||  | ||| | ||| ||||  | |
Db       2065 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2124

Qy       2212 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2271
                          || |    |   || ||||||  |||| || |   | | | |||    | |  | ||| |
Db       2125 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2184

Qy       2272 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2331
                          | |||||||| ||||| |||||||||||    |  |||| || |   | ||| |||||
Db       2185 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2244

Qy       2332 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2391
                          ||  | ||  |   |||||  | | |||| | |||| | ||      | |       |||
Db       2245 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2304

Qy       2392 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2451
                            ||    | |   |    |     | |    | ||     || |||| ||  |||| ||
Db       2305 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2364

Qy       2452 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2511
                          | |    |||||||  ||||||||| || |||| | |||  | |||  | ||    | ||
Db       2365 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2424

Qy       2512 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2571
                          | ||  | |||  |      ||    |||||||| ||  | || ||||||||   |  | |
Db       2425 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2484

Qy       2572 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2631
                          ||||   || |   |   | || || ||   |     ||||| | ||| | || | ||||||
Db       2485 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2544

Qy       2632 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2691
                          ||||| || || ||   |     | || | ||    |||||    || | | |
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Db   2545 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2604

Qy   2692 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2736
           | || | || || | |||| | | || || | ||
Db   2605 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2649

RESULT 12
US-09-969-532-15
; Sequence 15, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
; APPLICANT: Walke, D. Wade
; APPLICANT: Scoville, John
; TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
; FILE REFERENCE: LEX-0244-USA
; CURRENT APPLICATION NUMBER: US/09/969,532
; CURRENT FILING DATE: 2001-10-02
; PRIOR APPLICATION NUMBER: US 60/237,280
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 15
;   LENGTH: 2661
;   TYPE: DNA
;   ORGANISM: homo sapiens
US-09-969-532-15

  Query Match   16.0%; Score 439.8; DB 4; Length 2661;
  Best Local Similarity   51.2%; Pred. No. 9.9e-89;
  Matches 1313; Conservative   0; Mismatches 1147; Indels 105; Gaps   8;

Qy   172 CTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTG 231
           || || || ||| | | |||||| || |||| || || |||||| ||| || |
Db   157 CTGCCTCATTTCATAGAGGAGCCAGATGATGCTTATATTATCAAGAGCAACCCTATTGCA 216

Qy   232 CTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTG 291
           || |||||| || ||| |||| |||||| ||||||||| ||||||||| |||||||||
Db   217 CTCAGGTGCAAAGCGAGGCCAGCCATGCAGATATTCTTCAAATGCAACGGCGAGTGGGTC 276

Qy   292 CGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATG 351
           | ||| || || ||||| || | |||| |||| || || |
Db   277 CATCAGAACGAGCACGTCTCTGAAGAGACTCTGGACGAGAGCTCAGGTTTGAAGGTCCGC 336

Qy   352 GAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATAC 411
           || || ||| |||||| | |||||| ||||| ||| | | |||| ||||| ||
Db   337 GAAGTGTTCATCAATGTTACTAGGCAACAGGTGGAGGACTTCCATGGGCCCGAGGACTAT 396

Qy   412 TGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATC 471
           |||||||||||| ||||| |||||| | ||| ||| ||||||| |||||||| |
Db   397 TGGTGCCAGTGTGTGGCGTGGAGCCACCTGGGTACCTCCAAGAGCAGGAAGGCCTCTGTG 456

Qy   472 CGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTG 531
           ||||||||| || || || ||||| || || || || || | | | ||| || || |
Db   457 CGCATAGCCTATTTACGGAAAAACTTTGAACAAGACCCACAAGGAAGGGAAGTTCCCATT 516

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Qy    532  GAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAG 591
           ||       || || |||| |||||| |||||| |||||  ||||| | ||||||||||||
Db    517  GAAGGCATGATTGTACTGCACTGCCGCCCACCAGAGGGAGTCCCTGCTGCCGAGGTGGAA 576

Qy    592  TGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGG 651
           |||||    || || || |  | |||| |     | |||  || |  | ||| || |
Db    577  TGGCTGAAAAATGAAGAGCCCATTGACTCTGAACAAGACGAGAACATTGACACCAGGGCT 636

Qy    652  GAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTG 711
           ||  || |  |||| |   |  ||||||  || ||  |  || |   || ||||||||| ||
Db    637  GACCATAACCTGATCATCAGGCAGGCACGGCTCTCGGACTCAGGAAATTACACCTGCATG 696

Qy    712  GCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAAC 771
           ||    ||||||||||| |    |  |  ||  || || ||||    | ||||||||| |
Db    697  GCAGCCAACATCGTGGCTAAGAGGAGAAGCCTGTCGGCCACTGTTGTGGTCTACGTGGAT 756

Qy    772  GGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGG 831
           ||  | |||      ||||| |||  ||||||||||||||||| |      ||||
Db    757  GGGAGCTGGGAAGTGTGGAGCGAATGGTCCGTCTGCAGTCCAGAGTGTG---------AA 807

Qy    832  CAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAG 891
           ||    |||| |||| |     |||||   ||  | ||  || |||||||  ||||||||||
Db    808  CATTTGCGGATCCGGGAGTGCACAGCACCACCCCCGAGAAATGGGGGCAAATTCTGTGAA 867

Qy    892  GGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGC 951
           ||| | |    |       |     | ||||| |         |    ||||   ||
Db    868  GGTCTAAGCCAGGAATCTGAAAACTGCACAGATGGTCTTTGCATCCTAGGCATTGAGAAT 927

Qy    952  CCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGC 1011
            | |  || |  | || |  || ||  |||
Db    928  GCCAGCGACATTGCTTTGTACTCGGGCTTGG---------------------------- 958

Qy    1012 TCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGC 1071
                            |   |   |   | ||      || |  | |||| |      | |
Db    959  ----------------GTGCTGCCGTCGTGGCCGTTGCAGTCCTGGTCATTGGTGTCACC 1002

Qy    1072 AACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTAT 1131
              |  |  |   ||    |||  |  | |  |   | |     || |  |||     || |
Db    1003 CTTTACAGACGGAGCCAGAGTGACTATGGCGTGGACGTCATTGACTCTTCTGCATTGACA 1062

Qy    1132 GTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTT 1191
            |   ||| ||    || |   | ||    |||| |         ||| |||| |     |
Db    1063 GGTGGCTTCCAGACCTTCAACTTCAAAACAGTCCGTCAAGGTAACTCCCTGCTCCTGAAT 1122

Qy    1192 TATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCA 1251
            |..| | | |          ||| |  || | |   ||| |       |    .|.| ||
Db    1123 TCTGCCATGCA--------GCCAGATCTGACAGTGAGCCGGACATACAGCGGACCCATCT 1174

Qy    1252 GGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATC 1311
            | || |   | || || || || | |   | |||   || | | ||| || ||    | |
Db    1175 GTCTGCAGGACCCTCTGGACAAGG---AGCTCATGACAGAGTCCTCACTCTTTAACCCTT 1231

Qy    1312 CAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGAT 1371
            | ||      | |    |   |   |||   |  | |   || || ||    | | |||
Db    1232 TGTCGGACATCAAAGTGAAAGTCCAGAGCTCGTTCATGGTTTCCCTGGGAGTGTCTGAGA 1291

Qy    1372 GGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGC 1431
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                       | || ||  ||| |          |||       || |      |
Db   1292 GAGCTGAGTACCACGGCAAGAATCATTCCAGGACTTTT-------------------- 1329

Qy   1432 CGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTC 1491
          | |||   |    ||||||||| |   ||    |  ||    |   |  || | | ||
Db   1330 CCCCATGGAAACAACCACAGCTTTAGTACAATGCATCCCAGAAATAAAATGCCCTACATC 1389

Qy   1492 TCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACC 1551
           |   | |        || || |||  | |  ||      | |||    |||      |
Db   1390 CAAAATCTGT---------CATCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTC 1440

Qy   1552 TTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCC 1611
          ||  |   | ||||| ||   | | ||| || ||||||||| | ||| | ||||| ||
Db   1441 TTTGGCCATTTAGGGGGGCGCTTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCA 1500

Qy   1612 CCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAA 1671
          | | ||||||  ||    | |||    | ||||||| ||  |  | |   ||| | |
Db   1501 CACGGTGCCATCCCAGAGGAGAATTCTTGGGAGATTTATATGTCCAT---CAACCAAGGT 1557

Qy   1672 GACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGA 1731
          ||     || | |    || ||||| | ||    || |||||||| ||  | ||||||
Db   1558 GAACCCAGCCTCCAGTCAGATGGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGT 1617

Qy   1732 CCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCC 1791
           ||  ||  | |  | | |  |||||   || |   || | || |||||||  ||   |
Db   1618 CCTCCAGACATGATCGTCACCACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTC 1677

Qy   1792 AGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGAT 1851
          ||  ||||    ||||   || | | |  | ||  |  ||| |   |||||| |||||||
Db   1678 AGTTCTGAGCATTGGAATATCCATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAA 1737

Qy   1852 GTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGT 1911
          ||| ||   ||| || || ||  | | ||   ||  |||||||| |||| || |
Db   1738 GTGATGTCAGTGGAAGATGAATCTACATC------CTGTTACTGCCTTTTGGACCCCTTT 1791

Qy   1912 GCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGC 1971
          || ||  | ||  ||  ||  |  | || || || || || |||  |||||| |  |||
Db   1792 GCGTGTCATGTGCTCCTGGACAGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACA 1851

Qy   1972 GTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTC 2031
          |     ||||| |||| || || || ||| ||    | ||||    || |||| | ||||||
Db   1852 GACTGTGCCGTGAAGCAACTGAAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTG 1911

Qy   2032 GAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTG 2091
          || |||||| |  | || ||||| || |  | |||    |||| |    |||| |||||
Db   1912 GATTACAACTTGAGAGTTTACTGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTT.1971

Qy   2092 CAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAG 2151
          || | ||| | ||  ||  |||||||| |   | | || |||    | ||||| |||||
Db   1972 TCAGATGAAAGGCATCAAGGTGGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAA 2031

Qy   2152 GACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGT 2211
          | | | |  | | ||   |   || |  ||| ||| |    | |   | ||| ||||  | |
Db   2032 GGGAATACCTTTAGTCTTCAGATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATT 2091

Qy   2212 AAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGG 2271
          || |   |    || ||||||| |||| || |  | | |  |||       | |  | ||| |
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Db     2092 AAACCATTCACTGCCTGCCAGGAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAG 2151

Qy     2272 TACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGC 2331
            | ||||||||  ||||| |||||||||||  |  |||| || |   | ||| |||||
Db     2152 CCCCTGCACTGTGCCTTCTCCCTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGC 2211

Qy     2332 AAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATC 2391
            ||  | ||  |  ||||| |  | ||| | |  ||||  | ||    |  |       |||
Db     2212 AAAATCTGCATTCGGCAGCTCAAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATC 2271

Qy     2392 ACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCC 2451
            ||    | |    |      |     |         |      |   ||    |||| ||
Db     2272 CTAGAGAGTGAACGAGAAACCATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCA 2331

Qy     2452 CTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGC 2511
            | |   ||||||||  |||||||| || |||| |  ||| | ||| ||| | |  | |
Db     2332 CAGACTGGCCCCAAAGCCTTCAAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACA 2391

Qy     2512 CTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTG 2571
            | ||  | | |||  |      ||        ||||||| ||  | || ||||||  | | |
Db     2392 TTTGATACCCCCAATGCCAAAGGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATC 2451

Qy     2572 GACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTG 2631
            ||||  || |     |  || ||  | |    |||||  | ||  | ||  | ||||||
Db     2452 AACAGGAATTTATCTTATTTCGCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTG 2511

Qy     2632 TGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGA 2691
            |||||  ||  ||  ||     |   | ||  | ||    |  |||||    ||   |  |
Db     2512 TGGGAAGCTCGTCATCAGCATGATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAG 2571

Qy     2692 CTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAG 2736
            | ||   | | || || |    |||| |  | || || |  ||
Db     2572 ATTGGGAGGACACACACGAAACTCTCAAACATTTCAGAATCCCAG 2616


RESULT 13
US-09-471-276-345
; Sequence 345, Application US/09471276
; Patent No. 6822072
; GENERAL INFORMATION:
;  APPLICANT: Dumas Milne Edwards, J.B.
;  APPLICANT: Duclert A.
;  APPLICANT: Giordano, J.Y.
;  TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; Patent No. 6822072
;  FILE REFERENCE: GENSET.025CP1.
;  CURRENT APPLICATION NUMBER: US/09/471,276
;  CURRENT FILING DATE: 1999-12-21
;  EARLIER APPLICATION NUMBER: 09/057,719
;  EARLIER FILING DATE: 1998-04-09
;  EARLIER APPLICATION NUMBER: 09/069,047
;  EARLIER FILING DATE: 1998-04-28
;  EARLIER APPLICATION NUMBER: PCT/IB99/00712
;  EARLIER FILING DATE: 1999-04-09
;  NUMBER OF SEQ ID NOS: 1622
;  SOFTWARE: Patent.pm
; SEQ ID NO 345
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;   LENGTH: 349
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: 207..347
;   NAME/KEY: sig_peptide
;   LOCATION: 207..278
;   OTHER INFORMATION: Von Heijne matrix
;   OTHER INFORMATION: score 5.40000009536743
;   OTHER INFORMATION: seq SCCCLSSSSFIAG/RR
US-09-471-276-345
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   Best Local Similarity    98.8%;   Pred. No. 6.9e-63;
   Matches  335;  Conservative   1;  Mismatches   2;  Indels   1;  Gaps   1;
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Qy      934 GTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCAC 993
            || ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       12 GTGGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCAC 71

Qy      994 TGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGC 1053
            ||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       72 TGGCGGA-CCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGC 130

Qy     1054 ACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCT 1113
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Db      131 ACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACACTGCTTCTGGCCCT 190

Qy     1114 GAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTG 1173
            |||||||||||||||||||||||||||||||||||||||||||:||||||||||||||||
Db      191 GAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCBTGGTCCTGCTGCTG 250

Qy     1174 CTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGAC 1233
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      251 CTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGAC 310

Qy     1234 TCGTCCATTCTCACCTCAGGCTTCCAGCCCGTCAGCATC 1272
            |||||||||||||||||||||||||||||||||||||||
Db      311 TCGTCCATTCTCACCTCAGGCTTCCAGCCCGTCAGCATC 349
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```
RESULT 14
US-09-969-532-31
; Sequence 31, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
;  APPLICANT: Walke, D. Wade
;  APPLICANT:  Scoville, John
;  TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
;  FILE REFERENCE: LEX-0244-USA
;  CURRENT APPLICATION NUMBER: US/09/969,532
;  CURRENT FILING DATE:  2001-10-02
;  PRIOR APPLICATION NUMBER: US 60/237,280
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 33
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   Query Match              10.7%;  Score 293.2;  DB 4;  Length 1968;
   Best Local Similarity    53.7%;  Pred. No. 5.8e-56;
   Matches  657;  Conservative   0;  Mismatches  558;  Indels   9;  Gaps   2;


Qy      1513 TCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGG 1572
             II II III  I I II      I III   III   III  I   I IIIII II
Db       709 TCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTCTTTGGCCATTTAGGGGGGCGC 768


Qy      1573 CTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAGGG 1632
             I  I II II IIIIIIII I III I IIIII II I  I IIIIII II    I I
Db       769 TTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCACACGGTGCCATCCCAGAGGAG 828


Qy      1633 AAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCT 1692
             II    I  IIIII II I  I  I  II I   III  I  II  I I    II I
Db       829 AATTCTTGGGAGATTTATATGTCCATCAACCAAGGTGAACCC---AGCCTCCAGTCAGAT 885


Qy      1693 GGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACC 1752
             IIII I I II   II IIIIIIII    II I IIIIII II II I I I   IIIII
Db       886 GGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGTCCTCCAGACATGATCGTCACC 945


Qy      1753 CGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTG 1812
               II I   II I I II   IIIIIII II  III IIII    IIII    I
Db       946 ACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTCAGTTCTGAGCATTGGAATATC 1005


Qy      1813 CGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAG 1872
             I   I II III  I I      IIIIII IIIIIIII III II    III  II II
Db      1006 CATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAAGTGATGTCAGTGGAAGATGAA 1065


Qy      1873 GCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAG 1932
             I  I II    II IIIIIII  IIII II III II I II II   II
Db      1066 TCTACATC------CTGTTACTGCCTTTTGGACCCCTTTGCGTGTCATGTGCTCCTGGAC 1119


Qy      1933 CAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTC 1992
             I II   II III II  IIIIII I III I   IIIII IIII II
Db      1120 AGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACAGACTGTGCCGTGAAGCAACTG 1179


Qy      1993 AAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTAC 2052
             III II    I IIII     II IIII I IIIIII II IIIIII I  I II III
Db      1180 AAGGTGGCGGTTTTTGGCTGCATGTCCTGTAACTCCCTGGATTACAACTTGAGAGTTTAC 1239


Qy      2053 TGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAGAAGCAGCTGGGG 2112
             II  II I  I IIII     IIII I IIII IIIII       II I III I  II
Db      1240 TGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTTTCAGATGAAAGGCATCAAGGT 1299


Qy      2113 GGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTACCACAACCTGCGC 2172
             IIIIIIII I  I II III     I IIIII IIIII I I I  I   I   II I
Db      1300 GGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAAGGGAATACCTTTAGTCTTCAG 1359


Qy      2173 CTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTTGTCAGCTACCAG 2232
             I II  III  III I III   I III IIII I IIII I  I       II IIII
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Db      1360 ATTTCTGTCCTTGATATTCCCCCATTCCTCTGGAGAATTAAACCATTCACTGCCTGCCAG 1419

Qy      2233 GAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACC 2292
             ||  |||| || |   |  | | |||    ||  | || |  | |||||||  |||||  ||
Db      1420 GAAGTCCCGTTCTCCCGCGTGTGGTGCAGTAACCGGCAGCCCCTGCACTGTGCCTTCTCC 1479

Qy      2293 CTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTG 2352
             ||||||||||   |  |||| ||  |    |  ||| |||||||  | ||  |  ||||| |
Db      1480 CTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGCAAAATCTGCATTCGGCAGCTC 1539

Qy      2353 GAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGACACAAGGTTTGCT 2412
              | ||| || |   |||| | ||    |  |    |||   ||    | |        |
Db      1540 AAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATCCTAGAGAGTGAACGAGAAACC 1599

Qy      2413 GAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTC 2472
                      |         |     ||    |||| || | |   ||||||| ||||||
Db      1600 ATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCACAGACTGGCCCCAAAGCCTTC 1659

Qy      2473 AAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGCGG 2532
             || ||  ||||| |   |||   | |||   ||    | ||  | ||   | |||   |
Db      1660 AAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACATTTGATACCCCCAATGCCAAA 1719

Qy      2533 GGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCTTT 2592
             ||      |||||||| ||     | || ||||||| |   | |||| ||| |    | ||
Db      1720 GGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATCAACAGGAATTTATCTTATTTC 1779

Qy      2593 GCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCCCC 2652
             ||   |  |     |||||  | ||   | ||  | |||||||||||| || || ||    |
Db      1780 GCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTGTGGGAAGCTCGTCATCAGCAT 1839

Qy      2653 AACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTGGC 2712
              | || | ||  | |||||   ||  | |  |  | | || | ||| | ||
Db      1840 GATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAGATTGGGAGGACACACACGAAA 1899

Qy      2713 CTCTTCACAGTGTCGGAGGCTGAG 2736
             ||||  |   | || ||  |  ||
Db      1900 CTCTCAAACATTTCAGAATCCCAG 1923
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RESULT 15
US-09-969-532-29
; Sequence 29, Application US/09969532
; Patent No. 6777232
; GENERAL INFORMATION:
; APPLICANT: Walke, D. Wade
; APPLICANT: Scoville, John
; TITLE OF INVENTION: No. 6777232el Human Membrane Proteins and Polynucleotides
Encoding the Same
; FILE REFERENCE: LEX-0244-USA
; CURRENT APPLICATION NUMBER: US/09/969,532
; CURRENT FILING DATE: 2001-10-02
; PRIOR APPLICATION NUMBER: US 60/237,280
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 29
;   LENGTH: 2001

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   Query Match              10.7%;  Score 293.2;  DB 4;  Length 2001;
   Best Local Similarity    53.7%;  Pred. No. 5.8e-56;
   Matches  657;  Conservative    0;  Mismatches  558;  Indels    9;  Gaps    2;

Qy      1513 TCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTCCTCGGGGGCCGG 1572
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Db       742 TCACTCCCCACAAGGACAGAACTGAGGACAACTGGTGTCTTTGGCCATTTAGGGGGGCGC 801

Qy      1573 CTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCCATACCCCGAGGG 1632
             |  | || || |||||||||  | ||| | ||||| || |  | |||||| ||    | |
Db       802 TTAGTAATGCCAAATACAGGGGTGAGCTTACTCATACCACACGGTGCCATCCCAGAGGAG 861

Qy      1633 AAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGGTTGCCCCTAGCT 1692
             ||   |  |||||| ||  | | | || |   |||   |  || |  | |   || |
Db       862 AATTCTTGGGAGATTTATATGTCCATCAACCAAGGTGAACCC---AGCCTCCAGTCAGAT 918

Qy      1693 GGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGCGTCCTGCTCACC 1752
             |||| | ||   || |||||||||    || | ||||||| || || ||| | | |||||
Db       919 GGCTCTGAGGTGCTCCTGAGTCCTGAAGTCACCTGTGGTCCTCCAGACATGATCGTCACC 978

Qy      1753 CGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGACAGCTGGAGCCTG 1812
             ||  |   || | || |||||||  || ||| || | ||||  |||    |||   |
Db       979 ACTCCCTTTGCATTGACCATCCCGCACTGTGCAGATGTCAGTTCTGAGCATTGGAATATC 1038

Qy      1813 CGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCACCTGGGCGAGGAG 1872
             |  | || ||| | |   |||||| ||||||||| ||| || ||     ||| ||  ||
Db      1039 CATTTAAAGAAGAGGACACAGCAGGGCAAATGGGAGGAAGTGATGTCAGTGGAAGATGAA 1098

Qy      1873 GCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTACGTCTTCACCGAG 1932
             |  |  || || |||||||  ||| || |  ||| || ||| || | ||  ||    ||
Db      1099 TCTACATC------CTGTTACTGCCTTTTGGACCCCTTTGCGTGTCATGTGCTCCTGGAC 1152

Qy      1933 CAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCCGCCAAGCGCCTC 1992
             | ||   || ||| || ||  |||||| |   ||| |   |||   | ||||| |||| ||
Db      1153 AGCTTTGGGACCTATGCGCTCACTGGAGAGCCAATCACAGACTGTGCCGTGAAGCAACTG 1212

Qy      1993 AAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAACATCCGGGTCTAC 2052
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             ||   || |  | | |||       |||| |  | |||| ||||| |   |||  ||  ||
Db      1273 TGTGTGGACAATACCCCTTGTGCATTTCAGGAAGTGGTTTCAGATGAAAGGCATCAAGGT 1332

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Db      1333 GGACAGCTCCTGGAAGAACCAAAATTGCTGCATTTCAAAGGGAATACCTTTAGTCTTCAG 1392

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Qy      2233 GAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCACTGCACCTTCACC 2292
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Qy      2293 CTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGGGTGTGGCAGGTG 2352
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Db      1513 CTGGAGCGTTATACGCCCACTACCACCCAGCTGTCCTGCAAAATCTGCATTCGGCAGCTC 1572

Qy      2353 GAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGACACAAGGTTTGCT 2412
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Db      1573 AAAGGCCATGAACAGATCCTCCAAGTGCAGACATCAATCCTAGAGAGTGAACGAGAAACC 1632

Qy      2413 GAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGCCCCAGTGCCTTC 2472
                 |          |     ||      ` |||| || | |   |||||||  ||||||
Db      1633 ATCACTTTCTTCGCACAAGAGGACAGCACTTTCCCTGCACAGACTGGCCCCAAAGCCTTC 1692

Qy      2473 AAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCACCCTGTAGGCGG 2532
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Db      1693 AAAATTCCCTACTCCATCAGACAGCGGATTTGTGCTACATTTGATACCCCCAATGCCAAA 1752

Qy      2533 GGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCATCTCAGCTTCTTT 2592
             ||       ||||||| ||    | || |||||||   | | |  ||||  || |    | ||
Db      1753 GGCAAGGACTGGCAGATGTTAGCACAGAAAAACAGCATCAACAGGAATTTATCTTATTTC 1812

Qy      2593 GCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCGCGGCACTTCCCC 2652
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Db      1813 GCTACACAAAGTAGCCCATCTGCTGTCATTTTGAACCTGTGGGAAGCTCGTCATCAGCAT 1872

Qy      2653 AACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAGCCAGACGCTGGC 2712
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Db      1873 GATGGTGATCTTGACTCCCTGGCCTGTGCCCTTGAAGAGATTGGGAGGACACACACGAAA 1932

Qy      2713 CTCTTCACAGTGTCGGAGGCTGAG 2736
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OM nucleic - nucleic search, using sw model

Run on: March 6, 2005, 05:25:16 ; Search time 1488.37 Seconds
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Perfect score: 2752
Sequence: 1 ccgcggggccccgcgcccgg..........tgagtgctgaggccggccag 2752

Scoring table: IDENTITY_NUC
                 Gapop 10.0 , Gapext 1.0

Searched: 5401638 seqs, 2966923429 residues

Total number of hits satisfying chosen parameters: 10803276

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

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          Pred. No. is the number of results predicted by chance to have a
          score greater than or equal to the score of the result being printed,
          and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
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2	2752	100.0	2752	17	US-10-624-932-1	Sequence 1, Appli
3	2676.4	97.3	2881	10	US-09-970-944-1	Sequence 1, Appli
4	2356	85.6	3561	18	US-10-643-795A-77	Sequence 77, Appl
5	2356	85.6	3580	17	US-10-311-623-13	Sequence 13, Appl
6	2259	82.1	3014	10	US-09-933-261-1	Sequence 1, Appli
7	2259	82.1	3014	14	US-10-256-702-1	Sequence 1, Appli
8	2252.2	81.8	2697	16	US-10-240-154-15	Sequence 15, Appl
9	1562.4	56.8	1787	10	US-09-933-261-2	Sequence 2, Appli
10	1562.4	56.8	1787	14	US-10-256-702-2	Sequence 2, Appli
11	1206.6	43.8	1321	17	US-10-296-115-365	Sequence 365, App
12	936.2	34.0	2860	17	US-10-087-684-1	Sequence 1, Appli
13	936.2	34.0	2860	17	US-10-087-684-3	Sequence 3, Appli
14	936.2	34.0	2860	17	US-10-218-779-1	Sequence 1, Appli
15	936.2	34.0	2860	17	US-10-218-779-3	Sequence 3, Appli
16	913.6	33.2	2895	17	US-10-037-417-37	Sequence 37, Appl
17	904	32.8	3485	9	US-09-816-828-18	Sequence 18, Appl
18	902.4	32.8	3884	14	US-10-028-072-145	Sequence 145, App
19	902.4	32.8	3884	14	US-10-140-808-145	Sequence 145, App
20	902.4	32.8	3884	14	US-10-121-049-145	Sequence 145, App
21	902.4	32.8	3884	14	US-10-123-904-145	Sequence 145, App
22	902.4	32.8	3884	14	US-10-140-470-145	Sequence 145, App
23	902.4	32.8	3884	14	US-10-175-746-145	Sequence 145, App
24	902.4	32.8	3884	14	US-10-176-918-145	Sequence 145, App
25	902.4	32.8	3884	14	US-10-176-921-145	Sequence 145, App
26	902.4	32.8	3884	14	US-10-137-865-145	Sequence 145, App
27	902.4	32.8	3884	14	US-10-140-474-145	Sequence 145, App
28	902.4	32.8	3884	14	US-10-142-431-145	Sequence 145, App
29	902.4	32.8	3884	14	US-10-143-114-145	Sequence 145, App
30	902.4	32.8	3884	14	US-10-142-419-145	Sequence 145, App
31	902.4	32.8	3884	14	US-10-123-262-145	Sequence 145, App
32	902.4	32.8	3884	14	US-10-142-423-145	Sequence 145, App
33	902.4	32.8	3884	14	US-10-121-050-145	Sequence 145, App
34	902.4	32.8	3884	14	US-10-141-755-145	Sequence 145, App
35	902.4	32.8	3884	14	US-10-143-032-145	Sequence 145, App
36	902.4	32.8	3884	14	US-10-123-108-145	Sequence 145, App
37	902.4	32.8	3884	14	US-10-123-236-145	Sequence 145, App
38	902.4	32.8	3884	14	US-10-123-261-145	Sequence 145, App
39	902.4	32.8	3884	14	US-10-140-921-145	Sequence 145, App
40	902.4	32.8	3884	14	US-10-140-928-145	Sequence 145, App
41	902.4	32.8	3884	14	US-10-121-045-145	Sequence 145, App
42	902.4	32.8	3884	14	US-10-123-292-145	Sequence 145, App
43	902.4	32.8	3884	14	US-10-123-903-145	Sequence 145, App
44	902.4	32.8	3884	14	US-10-124-819-145	Sequence 145, App
45	902.4	32.8	3884	14	US-10-124-822-145	Sequence 145, App

ALIGNMENTS

RESULT 1
US-09-918-779-1

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Query Match            100.0%;  Score 2752;  DB 10;  Length 2752;
Best Local Similarity  100.0%;  Pred. No. 0;
Matches 2752;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;


Qy        1 CCGCGGGGCCCCGCGCCCGGCCCGCCCGCCTGCCCGCCCGCGGCCATGGCCGTCCGGCCC 60
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db        1 CCGCGGGGCCCCGCGCCCGGCCCGCCCGCCTGCCCGCCCGCGGCCATGGCCGTCCGGCCC 60


Qy       61 GGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTCCGCGGCTCGGGTGCC 120
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db       61 GGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTCCGCGGCTCGGGTGCC 120


Qy      121 CAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCAC 180
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      121 CAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCAC 180


Qy      181 TTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGC 240
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      181 TTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGC 240


Qy      241 AAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTG 300
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      241 AAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTG 300


Qy      301 GACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGC 360
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      301 GACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGC 360


Qy      361 ATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAG 420
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      361 ATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAG 420


Qy      421 TGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCC 480
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      421 TGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCC 480


Qy      481 AGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGC 540
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      481 AGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGC 540


Qy      541 ATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGG 600
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      541 ATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGG 600


Qy      601 AACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGC 660
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      601 AACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGC 660


Qy      661 CTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAAC 720
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      661 CTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAAC 720


Qy      721 ATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGG 780
            IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db      721 ATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGG 780
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Qy    781 TCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGG 840
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||.||
Db    781 TCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGG 840

Qy    841 AGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAAT 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    841 AGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAAT 900

Qy    901 GTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGC 960
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Db    901 GTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGC 960

Qy    961 AAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCA 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    961 AAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCA 1020

Qy   1021 GCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACC 1080
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1021 GCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACC 1080

Qy   1081 AGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTC 1140
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1081 AGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTC 1140

Qy   1141 ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGG 1200
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1141 ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGG 1200

Qy   1201 AAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTCCAG 1260
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1201 AAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTCCAG 1260

Qy   1261 CCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCGGAC 1320
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1261 CCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCGGAC 1320

Qy   1321 CTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATGGGCCCAGC 1380
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1321 CTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATGGGCCCAGC 1380

Qy   1381 CCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACACA 1440
          ||||||||.||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1381 CCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACACA 1440

Qy   1441 CTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAG 1500
          .  ||||||||||||||||||||||||||||||||||||||||||||||||.|||||||||||
Db   1441 CTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAG 1500

Qy   1501 AACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTC 1560
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1501 AACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTC 1560

Qy   1561 CTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCC 1620
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1561 CTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCC 1620

Qy   1621 ATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGG 1680
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                    |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1621 ATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGG 1680

Qy    1681 TTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGC 1740
                    |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1681 TTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGC 1740

Qy    1741 GTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGAC 1800
                    |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1741 GTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGAC 1800

Qy    1801 AGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCAC 1860
                    |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1801 AGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCAC 1860

Qy    1861 CTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTAC 1920
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Db    1861 CTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTAC 1920

Qy    1921 GTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCC 1980
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Db    1921 GTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCC 1980

Qy    1981 GCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAAC 2040
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Db    1981 GCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAAC 2040

Qy    2041 ATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAG 2100
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Db    2041 ATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAG 2100

Qy    2101 AAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTAC 2160
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Db    2101 AAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTAC 2160

Qy    2161 CACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTT 2220
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Db    2161 CACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTT 2220

Qy    2221 GTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCAC 2280
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Db    2221 GTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCAC 2280

Qy    2281 TGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGG 2340
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2281 TGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGG 2340

Qy    2341 GTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGAC 2400
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2341 GTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGAC 2400

Qy    2401 ACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGC 2460
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Db    2401 ACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGC 2460

Qy    2461 CCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCA 2520
                    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db      2461  CCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCA 2520

Qy      2521  CCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCAT 2580
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2521  CCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCAT 2580

Qy      2581  CTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCG 2640
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Db      2581  CTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCG 2640

Qy      2641  CGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAG 2700
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Db      2641  CGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAG 2700

Qy      2701  CCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGCCGGCCAG 2752
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Db      2701  CCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGCCGGCCAG 2752
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RESULT 2
US-10-624-932-1
; Sequence 1, Application US/10624932
; Publication No. US20040096877A1
; GENERAL INFORMATION:
; APPLICANT: Taupier, Raymond
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Rastelli, Luca
; APPLICANT: Spaderna, Steven
; APPLICANT: Shimkets, Richard
; APPLICANT: Zerhusen, Bryan
; APPLICANT: Spytek, Kimberly
; APPLICANT: Shenoy, Suresh
; APPLICANT: Li, Li
; APPLICANT: Gusev, Vladimir
; APPLICANT:· Grosse, William
; APPLICANT: Alsobrook, John
; APPLICANT: Lepley, Denise
; APPLICANT: Burgess, Catherine
; APPLICANT: Gerlach, Valerie
; APPLICANT: Ellerman, Karen
; APPLICANT: MacDougall, John
; APPLICANT: Stone, David
; APPLICANT: Smithson, Glennda
; TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-074 US
; CURRENT APPLICATION NUMBER: US/10/624,932
; CURRENT FILING DATE: 2003-07-21
; PRIOR APPLICATION NUMBER: 09/918,779
; PRIOR FILING DATE: 2001-07-03
; PRIOR APPLICATION NUMBER: 60/221,409
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: 60/222,840
; PRIOR FILING DATE: 2000-08-04
; PRIOR APPLICATION NUMBER: 60/223,752
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/223,762
; PRIOR FILING DATE: 2000-08-08

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  Query Match                100.0%;  Score 2752;  DB 17;  Length 2752;
  Best Local Similarity    100.0%;  Pred. No. 0;
  Matches 2752;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;


Qy         1 CCGCGGGGCCCCGCGCCCGGCCCGCCCGCCTGCCCGCCCGCGGCCATGGCCGTCCGGCCC 60
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         1 CCGCGGGGCCCCGCGCCCGGCCCGCCCGCCTGCCCGCCCGCGGCCATGGCCGTCCGGCCC 60


Qy        61 GGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTCCGCGGCTCGGGTGCC 120
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Db        61 GGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTCCGCGGCTCGGGTGCC 120


Qy       121 CAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCAC 180
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       121 CAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCAC 180


Qy       181 TTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGC 240
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       181 TTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGC 240


Qy       241 AAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTG 300
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       241 AAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTG 300


Qy       301 GACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGC 360
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       301 GACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGC 360


Qy       361 ATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAG 420
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       361 ATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAG 420


Qy       421 TGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCC 480
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Db       421 TGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCC 480


Qy       481 AGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGC 540
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Db    481 AGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGC 540

Qy    541 ATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGG 600
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Db    541 ATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGG 600

Qy    601 AACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGC 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    601 AACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGC 660

Qy    661 CTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAAC 720
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Db    661 CTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAAC 720

Qy    721 ATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGG 780
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Db    721 ATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGG 780

Qy    781 TCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGG 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    781 TCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGG 840

Qy    841 AGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAAT 900
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Db    841 AGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAAT 900

Qy    901 GTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGC 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    901 GTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGC 960

Qy    961 AAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCA 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    961 AAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCA 1020

Qy   1021 GCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACC 1080
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1021 GCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACC 1080

Qy   1081 AGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTC 1140
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Db   1081 AGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTC 1140

Qy   1141 ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGG 1200
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Db   1141 ATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGG 1200

Qy   1201 AAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTCCAG 1260
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1201 AAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTCCAG 1260

Qy   1261 CCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCGGAC 1320
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1261 CCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCGGAC 1320

Qy   1321 CTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATGGGCCCAGC 1380
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1321 CTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATGGGCCCAGC 1380
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Qy      1381 CCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACACA 1440
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Db      1381 CCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACACA 1440

Qy      1441 CTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAG 1500
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Db      1441 CTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAG 1500

Qy      1501 AACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAACTTC 1560
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Qy      1561 CTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCC 1620
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Db      1561 CTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGATGCC 1620

Qy      1621 ATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTGAGG 1680
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Qy      1681 TTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGC 1740
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Db      1681 TTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCTGGC 1740

Qy      1741 GTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGAC 1800
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Db      1741 GTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCTGAC 1800

Qy      1801 AGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCAC 1860
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Db      1801 AGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATGTGCTGCAC 1860

Qy      1861 CTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTAC 1920
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Db      1861 CTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCCTGCTAC 1920

Qy      1921 GTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCC 1980
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Db      1921 GTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTGGCTGCC 1980

Qy      1981 GCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAAC 2040
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Db      1981 GCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAGTACAAC 2040

Qy      2041 ATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAG 2100
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2041 ATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAGCTGGAG 2100

Qy      2101 AAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTAC 2160
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Db      2101 AAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGACAGTTAC 2160

Qy      2161 CACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAGCTCCTT 2220
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy      2221 GTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCAC 2280
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Db      2221 GTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTACTTGCAC 2280

Qy      2281 TGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGG 2340
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Db      2281 TGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAGCTGTGG 2340

Qy      2341 GTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGAC 2400
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Db      2341 GTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACCAAGGAC 2400

Qy      2401 ACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGC 2460
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Db      2401 ACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTGGTGGGC 2460

Qy      2461 CCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCA 2520
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2461 CCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTGGACCCA 2520

Qy      2521 CCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCAT 2580
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Db      2521 CCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGACAGCCAT 2580

Qy      2581 CTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCG 2640
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Db      2581 CTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGGGAGGCG 2640

Qy      2641 CGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAG 2700
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Db      2641 CGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTGGGCCAG 2700

Qy      2701 CCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGCCGGCCAG 2752
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RESULT 3
US-09-970-944-1
; Sequence 1, Application US/09970944
; Publication No. US20030204052A1
; GENERAL INFORMATION:
;   APPLICANT: Herrman, John L
;   APPLICANT: Rastelli, Luca
;   APPLICANT: Shimkets, Richard A
;   TITLE OF INVENTION: No. US20030204052A1e1 Proteins and Nucleic Acids Encoding
Same and
;   TITLE OF INVENTION: Antibodies Directed Against these Proteins
;   FILE REFERENCE: 21402-138
;   CURRENT APPLICATION NUMBER: US/09/970,944
;   CURRENT FILING DATE: 2002-05-02
;   PRIOR APPLICATION NUMBER: 60/237,862
;   PRIOR FILING DATE: 2000-10-04
;   NUMBER OF SEQ ID NOS: 62
;   SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
;   LENGTH: 2881

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;    TYPE: DNA
;    ORGANISM: Homo sapiens
US-09-970-944-1

   Query Match          97.3%;   Score 2676.4;   DB 10;   Length 2881;
   Best Local Similarity  98.9%;   Pred. No. 0;
   Matches 2728;  Conservative   0;  Mismatches   21;  Indels   9;  Gaps   3;

Qy        1 CCGCGGGGCCCCGCGCCCGGCCCGCCCGCCTGCCCGCCCGCGGCCATGGCCGTCCGGCCC 60
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       42 CCGCGGGGCCCCGCGCCCGGCCCGCCCGCCTGCCCGCCCGCGGCCATGGCCGTCCGGCCC 101

Qy       61 GGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTCCGCGGCTCGGGTGCC 120
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Db      102 GGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTCCGCGGCTCGGGTGCC 161

Qy      121 CAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCAC 180
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Db      162 CAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCAC 221

Qy      181 TTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGC 240
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Db      222 TTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGC 281

Qy      241 AAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTG 300
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Db      282 AAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTG 341

Qy      301 GACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGC 360
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Db      342 GACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGTGAGCCGACCATGGAGGTCCGC 401

Qy      361 ATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAG 420
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Db      402 ATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAG 461

Qy      421 TGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCC 480
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Db      462 TGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCC 521

Qy      481 AGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGC 540
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Qy      541 ATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGG 600
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Db      582 ATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGG 641

Qy      601 AACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGC 660
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      642 AACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATCACGCGGGAGCACAGC 701

Qy      661 CTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAAC 720
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Db      702 CTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAAC 761

Qy      721 ATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGG 780
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Db      762 ATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGG 821

Qy      781 TCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGG 840
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Db      822 TCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGG 881

Qy      841 AGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAAT 900
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Db      882 AGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAAT 941

Qy      901 GTCCAGAA---AACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGG 957
            |||||  |   || | ||| |     ||   | | || ||||||||||||||||||||||
Db      942 GTCCATGACCGCACCGTCTCCTCTCTGCTTGTCTCTGTGGACGGCAGCTGGAGCCCGTGG 1001

Qy      958 AGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGAC 1017
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Db     1002 AGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGAC 1061

Qy     1018 CCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGT 1077
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Db     1062 CCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGACACCCGCAACTGT 1121

Qy     1078 ACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGC 1137
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1122 ACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGC 1181

Qy     1138 CTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGC 1197
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Db     1182 CTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGC 1241

Qy     1198 CGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTC 1257
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1242 CGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCTCAGGCTTC 1301

Qy     1258 CAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCG 1317
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1302 CAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACCATCCAGCCG 1361

Qy     1318 GACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATGGGCCC 1377
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Db     1362 GACCTCAG---CACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGATGGGCCC 1418

Qy     1378 AGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCAC 1437
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1419 AGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCAC 1478

Qy     1438 ACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACC 1497
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Db     1479 ACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACC 1538

Qy     1498 CAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAAC 1557
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1539 CAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTCAAC 1598

Qy     1558 TTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGAT 1617
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db    1599 TTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCAGAT 1658

Qy    1618 GCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTG 1677
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Db    1659 GCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGACGTG 1718

Qy    1678 AGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCT 1737
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Db    1719 AGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCCCCT 1778

Qy    1738 GGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCAGCCCT 1797
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Qy    1798 GACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGG---AGGATGTG 1854
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Qy    1855 CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCC 1914
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Db    1899 CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTGCC 1958

Qy    1915 TGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCGTG 1974
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Qy    1975 GCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAG 2034
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Db    2019 GCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCGAG 2078

Qy    2035 TACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAG 2094
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Db    2079 TACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGCAG 2138

Qy    2095 CTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGAC 2154
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Db    2139 CTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGGAC 2198

Qy    2155 AGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAG 2214
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2199 AGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTAAG 2258

Qy    2215 CTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTAC 2274
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Db    2259 CTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGTAC 2318

Qy    2275 TTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAG 2334
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Db    2319 TTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCAAG 2378

Qy    2335 CTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACC 2394
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Db    2379 CTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCACC 2438

Qy    2395 AAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTG 2454
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Db    2439 AAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGCCCTG 2498
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Qy    .  2455 GTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTG 2514
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Db       2499 GTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAGCCTG 2558

Qy       2515 GACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGAC 2574
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Db       2559 GACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCTGGAC 2618

Qy       2575 AGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGG 2634
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2619 AGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCTGTGG 2678

Qy       2635 GAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTG 2694
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2679 GAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGGACTG 2738

Qy       2695 GGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGCCGGCCAG 2752
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2739 GGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGCCGGCCAG 2796
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RESULT 4
US-10-643-795A-77
; Sequence 77, Application US/10643795A
; Publication No. US20040241703A1
; GENERAL INFORMATION:
;   APPLICANT: FREDERIC J. DESAUVAGE
;   APPLICANT: GRETCHEN FRANTZ
;   APPLICANT: KENNETH J. HILLAN
;   APPLICANT: PAUL POLAKIS
;   APPLICANT: ANDREW POLSON
;   APPLICANT: VICTORIA SMITH
;   APPLICANT: SUSAN D. SPENCER
;   APPLICANT: THOMAS D. WU
;   APPLICANT: ZEMIN ZHANG
;   TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
;   TITLE OF INVENTION: TREATMENT OF TUMOR
;   FILE REFERENCE: P5026R1-US
;   CURRENT APPLICATION NUMBER: US/10/643,795A
;   CURRENT FILING DATE: 2003-08-19
;   PRIOR APPLICATION NUMBER: US 60/404,809
;   PRIOR FILING DATE: 2002-08-19
;   PRIOR APPLICATION NUMBER: US 60/405,645
;   PRIOR FILING DATE: 2002-08-21
;   PRIOR APPLICATION NUMBER: US 60/413,192
;   PRIOR FILING DATE: 2002-09-23
;   PRIOR APPLICATION NUMBER: US 60/419,008
;   PRIOR FILING DATE: 2002-10-15
;   PRIOR APPLICATION NUMBER: US 60/426,847
;   PRIOR FILING DATE: 2002-11-15
;   PRIOR APPLICATION NUMBER: US 60/484,959
;   PRIOR FILING DATE: 2003-07-02
;   NUMBER OF SEQ ID NOS: 158
; SEQ ID NO 77
;   LENGTH: 3561
;   TYPE: DNA

Query Match   85.6%;   Score 2356;   DB 18;   Length 3561;
Best Local Similarity   93.6%;   Pred. No. 0;
Matches 2537;   Conservative   0;   Mismatches   5;   Indels   168;   Gaps   1;

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Qy      43 GCCATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGG 102
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1 GCCATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGG 60

Qy     103 CTCCGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAAC 162
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      61 CTCCGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAAC 120

Qy     163 CCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAG 222
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     121 CCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAG 180

Qy     223 CCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGG 282
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     181 CCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGG 240

Qy     283 GAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTG 342
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     241 GAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTG 300

Qy     343 CCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTG 402
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     301 CCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTG 360

Qy     403 GAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAG 462
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     361 GAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAG 420

Qy     463 GCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAG 522
           ||||||||||||||||||    ||||||||||||||||||||||||||||||||||||||
Db     421 GCCTACATCCGCATAGCCTATTTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAG 480

Qy     523 GTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCC 582
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     481 GTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCC 540

Qy     583 GAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATAC 642
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     541 GAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATAC 600

Qy     643 ATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTAC 702
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     601 ATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTAC 660

Qy     703 ACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTC 762
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     661 ACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTC 720

Qy     763 TACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGG 822
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Db      721  TACGTG------------------------------------------------------- 726

Qy      823  CGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCT 882

Db      727  ------------------------------------------------------------ 726

Qy      883  TTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGC 942
                                                                        ||||||
Db      727  ------------------------------------------------------GACGGC 732

Qy      943  AGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGC 1002
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      733  AGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGC 792

Qy      1003 CGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTG 1062
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      793  CGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTG 852

Qy      1063 GACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTG 1122
             |||||||||||||||||||||||||||||||||||||||| |||||||||||||||||||||
Db      853  GACACCCGCAACTGTACCAGTGACCTCTGTGTACACACTGCTTCTGGCCCTGAGGACGTG 912

Qy      1123 GCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTC 1182
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      913  GCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTC 972

Qy      1183 ATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT 1242
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      973  ATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT 1032

Qy      1243 CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTG 1302
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1033 CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTG 1092

Qy      1303 CTCACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCC 1362
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1093 CTCACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCC 1152

Qy      1363 CGGCAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTG 1422
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1153 CGGCAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTG 1212

Qy      1423 GGTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTC 1482
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1213 GGTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTC 1272

Qy      1483 TCCCGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACC 1542
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1273 TCCCGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACC 1332

Qy      1543 TATGGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTC 1602
             |||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||||
Db      1333 TATGGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTC 1392

Qy      1603 CTCATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCAC 1662
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1393 CTCATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCAC 1452
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Qy    1663 AAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTT 1722
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1453 AAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTT 1512

Qy    1723 AGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGT 1782
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1513 AGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGT 1572

Qy    1783 GGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGC 1842
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1573 GGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGC 1632

Qy    1843 TGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTG 1902
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1633 TGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTG 1692

Qy    1903 GAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAG 1962
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1693 GAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAG 1752

Qy    1963 GCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGC 2022
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1753 GCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGC 1812

Qy    2023 ACCTCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAG 2082
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1813 ACCTCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAG 1872

Qy    2083 GAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTG 2142
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1873 GAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTG 1932

Qy    2143 CACTTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTG 2202
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1933 CACTTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTG 1992

Qy    2203 TGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGC 2262
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1993 TGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGC 2052

Qy    2263 ACGCAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGAC 2322
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2053 ACGCAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGAC 2112

Qy    2323 CTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAAC 2382
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2113 CTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAAC 2172

Qy    2383 TTCAACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGG 2442
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2173 TTCAACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGG 2232

Qy    2443 GTCCCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATA 2502
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    2233 GTCCCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATA 2292
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Qy      2503 ATTTCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAA 2562
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2293 ATTTCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAA 2352

Qy      2563 CTCCACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATC 2622
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2353 CTCCACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATC 2412

Qy      2623 CTCAACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCA 2682
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2413 CTCAACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCA 2472

Qy      2683 GTGGCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2473 GTGGCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2532

Qy      2743 GGCCGGCCAG 2752
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Db      2533 GGCCGGCCAG 2542


RESULT 5
US-10-311-623-13
; Sequence 13, Application US/10311623
; Publication No. US20040023244A1
; GENERAL INFORMATION:
;  APPLICANT: INCYTE GENOMICS, INC.; GRIFFIN, Jennifer A.
;  APPLICANT:  KALLICK, Deborah A.; TRIBOULEY, Catherine M.
;  APPLICANT:  YUE, Henry; NGUYEN, Danniel B.
;  APPLICANT:  TANG, Y. Tom; LAL, Preeti G.
;  APPLICANT:  POLICKY, Jennifer L.; AZIMZAI, Yalda
;  APPLICANT:  LU, Dyung Aina M.; GRAUL, Richard C.
;  APPLICANT:  YAO, Monique G.; BURFORD, Neil
;  APPLICANT:  HAFALIA, April J. A.; BAUGHN, Mariah R.
;  APPLICANT:  BANDMAN, Olga; ARVIZU, Chandra S.
;  APPLICANT:  YANG, Junming; XU, Yuming
;  APPLICANT:  GANDHI, Ameena R.; WARREN, Bridget A.
;  APPLICANT:  DING, Li; SANJANWALA, Madhusudan M.
;  APPLICANT:  DUGGAN, Brendan M.; LU, Yan
;  TITLE OF INVENTION: RECEPTORS
;  FILE REFERENCE: PF-0793 USN
;  CURRENT APPLICATION NUMBER: US/10/311,623
;  CURRENT FILING DATE:  2002-12-17
;  PRIOR APPLICATION NUMBER: US 01/19942
;  PRIOR FILING DATE: 2001-06-21
;  PRIOR APPLICATION NUMBER: US 60/214,027
;  PRIOR FILING DATE: 2000-06-21
;  PRIOR APPLICATION NUMBER: US 60/228,045
;  PRIOR FILING DATE: 2000-08-25
;  PRIOR APPLICATION NUMBER: US 60/255,104
;  PRIOR FILING DATE: 2000-12-12
;  NUMBER OF SEQ ID NOS: 24
;  SOFTWARE: PERL Program
; SEQ ID NO 13
;   LENGTH: 3580
;   TYPE: DNA
;   ORGANISM: Homo sapiens
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   Query Match              85.6%;  Score 2356;  DB 17;  Length 3580;
   Best Local Similarity    93.6%;  Pred. No. 0;
   Matches 2537;  Conservative   0;  Mismatches    5;  Indels  168;  Gaps    1;

Qy       43 GCCATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGG 102
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1 GCCATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGG 60

Qy      103 CTCCGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAAC 162
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       61 CTCCGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAAC 120

Qy      163 CCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAG 222
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      121 CCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAG 180

Qy      223 CCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGG 282
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Db      181 CCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGG 240

Qy      283 GAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTG 342
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      241 GAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTG 300

Qy      343 CCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTG 402
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      301 CCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTG 360

Qy      403 GAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAG 462
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      361 GAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAG 420

Qy      463 GCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAG 522
            |||||||||||||||||      ||||||||||||||||||||||||||||||||||||||
Db      421 GCCTACATCCGCATAGCCTATTTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAG 480

Qy      523 GTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCC 582
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Db      481 GTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCC 540

Qy      583 GAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATAC 642
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      541 GAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATAC 600

Qy      643 ATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTAC 702
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      601 ATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTAC 660

Qy      703 ACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTC 762
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      661 ACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTC 720
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Qy    763  TACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGG 822
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Db    721  TACGTG------------------------------------------------------ 726


Qy    823  CGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCT 882

Db    727  ------------------------------------------------------------ 726


Qy    883  TTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGC 942
                                                                  ||||||
Db    727  ---------------------------------------------------GACGGC 732


Qy    943  AGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGC 1002
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    733  AGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGC 792


Qy    1003 CGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTG 1062
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    793  CGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTG 852


Qy    1063 GACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTG 1122
           ||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||||
Db    853  GACACCCGCAACTGTACCAGTGACCTCTGTGTACACACTGCTTCTGGCCCTGAGGACGTG 912


Qy    1123 GCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTC 1182
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    913  GCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTC 972


Qy    1183 ATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT 1242
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    973  ATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATT 1032


Qy    1243 CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTG 1302
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1033 CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTG 1092


Qy    1303 CTCACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCC 1362
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1093 CTCACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCC 1152


Qy    1363 CGGCAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTG 1422
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1153 CGGCAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTG 1212


Qy    1423 GGTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTC 1482
           |||||||||||||||||||||||||||||||||.||||||||||||||||||||||||||
Db    1213 GGTGGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTC 1272


Qy    1483 TCCCGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACC 1542
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1273 TCCCGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACC 1332


Qy    1543 TATGGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTC 1602
           |||||||||||||||||||||||||||||||||||||||||||||||||| |||||||||
Db    1333 TATGGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTC 1392


Qy    1603 CTCATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCAC 1662
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Db      1393 CTCATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCAC 1452

Qy      1663 AAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTT 1722
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1453 AAGCCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTT 1512

Qy      1723 AGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGT 1782
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1513 AGCTGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGT 1572

Qy      1783 GGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGC 1842
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Db      1573 GGGGAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGC 1632

Qy      1843 TGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTG 1902
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1633 TGGGAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTG 1692

Qy     ·1903 GAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAG 1962
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1693 GAGGCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAG 1752

Qy      1963 GCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGC 2022
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1753 GCCCTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGC 1812

Qy      2023 ACCTCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAG 2082
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1813 ACCTCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAG 1872

Qy      2083 GAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTG 2142
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1873 GAGGTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTG 1932

Qy      2143 CACTTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTG 2202
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1933 CACTTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTG 1992

Qy      2203 TGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGC 2262
                        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1993 TGGAAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGC 2052

Qy      2263 ACGCAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGAC 2322
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2053 ACGCAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGAC 2112

Qy      2323 CTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAAC 2382
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2113 CTGGCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAAC 2172

Qy      2383 TTCAACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGG 2442
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2173 TTCAACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGG 2232

Qy      2443 GTCCCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATA 2502
                        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        2233 GTCCCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATA 2292

Qy        2503 ATTTCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAA 2562
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        2293 ATTTCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAA 2352

Qy        2563 CTCCACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATC 2622
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        2353 CTCCACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATC 2412

Qy        2623 CTCAACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCA 2682
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Db        2413 CTCAACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCA 2472

Qy        2683 GTGGCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        2473 GTGGCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2532

Qy        2743 GGCCGGCCAG 2752
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Db        2533 GGCCGGCCAG 2542


RESULT 6
US-09-933-261-1
; Sequence 1, Application US/09933261
; Publication No. US20030040046A1
;   GENERAL INFORMATION:
;         APPLICANT: Tessier-Lavigne, Marc
;                    Leonardo, E. David
;                    Hink, Lindsay
;                    Masu, Masayuki
;                    Kazuko, Keino-Masu
;         TITLE OF INVENTION: Netrin Receptors
;         NUMBER OF SEQUENCES: 8
;         CORRESPONDENCE ADDRESS:
;             ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;             STREET: 268 BUSH STREET, SUITE 3200
;             CITY: SAN FRANCISCO
;             STATE: CALIFORNIA
;             COUNTRY: USA
;             ZIP: 94104
;         COMPUTER READABLE FORM:
;             MEDIUM TYPE: Floppy disk
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: PatentIn Release #1.0, Version #1.30
;         CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/933,261
;             FILING DATE: 20-Aug-2001
;             CLASSIFICATION: <Unknown>
;         PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 08/808,982
;             FILING DATE: <Unknown>
;         ATTORNEY/AGENT INFORMATION:
;             NAME: OSMAN, RICHARD A
;             REGISTRATION NUMBER: 36,627
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  Query Match          82.1%;  Score 2259;  DB 10;  Length 3014;
  Best Local Similarity  89.7%;  Pred. No. 0;
  Matches 2427;  Conservative   0;  Mismatches  280;  Indels   0;  Gaps    0;
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Qy        46 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 105
             ||||||||||||||||||||||||||||||||| |||||||||||||||||||||| |||||
Db         1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGTGCTCCTGGGCATAGTCCTCGCCGCCTGGCTT 60

Qy       106 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 165
             || || ||||||||||||||||||||||||| ||||||||| ||||||||| |||||||||||
Db        61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120

Qy       166 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 225
             |||||||| |||||||||||||| ||||| ||||| ||||||||| |||||||||||||||||
Db       121 GACCTGCTGCCCCACTTCCTGGTAGAGCCTGAGGACGTGTACATTGTCAAGAACAAGCCG 180

Qy       226 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 285
             ||| || | ||||||||||| ||||| ||||| ||||||||||||||||||||||| |||||
Db       181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240

Qy       286 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 345
             ||||| |||||||| || |||||| || || ||||||||| ||| | |||||| || |||||
Db       241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300

Qy       346 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 405
             |||||||||||||| || || || || |||||||||||||| ||||| ||||| |||||||||
Db       301 ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG 360

Qy       406 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 465
             ||||||||||||||||| |||||||||||||||||||| ||||||||| ||||||||||||||
Db       361 GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC 420

Qy       466 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 525
             |||||||| || ||||    ||||||||||||| |||||||||||| |||||||||||| |||
Db       421 TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG 480

Qy       526 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 585
             || ||||||||| ||||| || || || || || || || || || || ||||| ||||| |||
Db       481 TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCCAGAAGGAATCCCCCCAGCTGAG 540

Qy       586 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 645
             ||||||||||| || || ||||||||| ||||||||| ||||| || ||||||||| ||||||
Db       541 GTGGAGTGGCTTCGAAATGAGGACCTCGTGGACCCCTCCCTCGATCCCAATGTGTACATC 600
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Qy     646 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 705
           |||||||||||||||| || |||||| |||||||||||| || |||||||||||||||||
Db     601 ACGCGGGAGCACAGCCTAGTCGTGCGTCAGGCCCGCCTGGCCGACACGGCCAACTACACC 660

Qy     706 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 765
           || ||||||||||||||||||| || |||||||| ||| |||| || || |||||| || ||
Db     661 TGTGTGGCCAAGAACATCGTAGCCCGTCGCCGAAGCACCTCTGCAGCGGTCATTGTTTAT 720

Qy     766 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 825
           |||||||||||||||||||||||||||| |||||||||||||||||||||||||||||||
Db     721 GTGAACGGTGGGTGGTCGACGTGGACTGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGT 780

Qy     826 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 885
           |||||||||||||||||||||||||||||||||||||||||| ||||||||||||| |||
Db     781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC 840

Qy     886 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 945
           ||||||||||||||||||||||||||||||||||||||||| ||||||||| || || |||
Db     841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACTCTGTGCCCAGTGGATGGGAGC 900

Qy     946 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 1005
           |||||    ||||||| |||||||| |||||||| ||||||||||||||||||||||||||
Db     901 TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC 960

Qy    1006 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1065
           |||||||||||||||||||||||||| ||||| ||||||| | ||| ||||||||||||
Db     961 GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC 1020

Qy    1066 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1125
           |||||||||||||||||||||||||| | |||| ||||||| |||| ||||||||||
Db    1021 ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT 1080

Qy    1126 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1185
           |||||  | ||||| |||| ||||  |||| || |||||  || ||||| ||||| | |||
Db    1081 CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA 1140

Qy    1186 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1245
           ||| |||| || || ||||||||| ||||||||||| ||||||||| ||||||||||| |||
Db    1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC 1200

Qy    1246 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1305
           ||||| |||||||||||| ||||||||||||||||||||||||||||||||||| ||||||
Db    1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC 1260

Qy    1306 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1365
           |||||||||| ||||||||||||||||||| |||||||||||||||||||| ||| | ||
Db    1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG 1320

Qy    1366 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1425
           |||||||| ||||||||||||||||||||| | ||||| ||||||||||||||||| |||||
Db    1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG 1380

Qy    1426 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1485
           | |||||||| || ||||||||||||||||| |||||||||||||| ||||| |||||||||
Db    1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC 1440
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Qy    1486 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1545
           .|||||||||||||| |||||||| || ||||||||||| ||||||||||||||||| ||||
Db    1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC 1500

Qy    1546 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1605
           ||||||||||||||||||||||||||||||||||||||||||||||||| || |||||||||||
Db    1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC 1560

Qy    1606 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1665
           || |||||| |||||||| |||||||| |||||||| ||||||||||||||||| |||||||||
Db    1561 ATACCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG 1620

Qy    1666 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1725
           || ||||||||||||||||||||||||||||||||||||||||||||||||||   |||||||||
Db    1621 CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC 1680

Qy    1726 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1785
           ||||| ||||| || ||||||||||||||||||||||||||||||| || |||||||||||||||
Db    1681 TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA 1740

Qy    1786 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1845
           |||||||||||||||||||||||||| |||||||||||||||||||||| || |||||||||||| |||
Db    1741 GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG 1800

Qy    1846 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1905
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Db    1801 GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG 1860

Qy    1906 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1965
           ||| | ||||||||| ||||||||| |||||||||||||||||||||||||| |||||||||
Db    1861 GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC 1920

Qy    1966 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 2025
           |||||||||||||| |||||||||||||| ||| ||||||||||| || |||||||| ||
Db    1921 CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG 1980

Qy    2026 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2085
           ||||| ||||||||||||||||| || |||||||| || ||||||||||| || |||||||||
Db    1981 TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG 2040

Qy    2086 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2145
           ||||||||||||||||||||||||| || |||||||||||||||||||||| || |||||||||
Db    2041 GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC 2100

Qy    2146 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2205
           |||||| ||||||||||||||||||| ||..|| |||||||||| ||||||||||||||||||| 
Db    2101 TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160

Qy    2206 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2265
           ||||| ||||| ||||||||||||||||||||||||| ||||| ||||||||||| |||||
Db    2161 AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy    2266 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2325
           |||| ||| |||||||||||||||||||||||||||| ||| | |||||||| || ||||||
Db    2221 CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280

Qy    2326 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2385
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            ||||||||||  |||||||||||||||||||||||||  ||  |||||||||||||  |||||||||||
Db    2281  GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy    2386  AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2445
            ||||||||||  ||||||||||||||||||||||||  ||  |||||||||||||||||||  |||||||
Db    2341  AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy    2446  CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2505
            |||||||||||||||||||||||||||||||||||||||||||||||||||||||  |||||  ||
Db    2401  CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy    2506  TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2565
             ||||  ||||||||||||||  ||  ||||||  |||||||||  |  |||||  ||||||||||||
Db    2461  GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy    2566  CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2625
            |||||||||||||||||||||  |||||||||||||||||||||||||||  |||||||||||||||||
Db    2521  CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy    2626  AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2685
            |||||  |||||||||  |||||||||||||||||||||||||||  ||||||||||  |||||  |||
Db    2581  AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy    2686  GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGC 2745
            ||  |||||||||||||  |||||  |||||||||||||||  ||||||||||  |||||  ||||  |
Db    2641  GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGAGAC 2700

Qy    2746  CGGCCAG 2752
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Db    2701  CAGCCAG 2707
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RESULT 7
US-10-256-702-1
; Sequence 1, Application US/10256702
; Publication No. US20030059859A1
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay
;   Masu, Masayuki
;   Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 8
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;   STREET: 268 BUSH STREET, SUITE 3200
;   CITY: SAN FRANCISCO
;   STATE: CALIFORNIA
;   COUNTRY: USA
;   ZIP: 94104
;   COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:

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   Query Match           82.1%;  Score 2259;  DB 14;  Length 3014;
   Best Local Similarity  89.7%;  Pred. No. 0;
   Matches 2427;  Conservative  0;  Mismatches  280;  Indels  0;  Gaps  0;


Qy      46 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 105
           ||||||||||||||||||||||||||||||| ||||||||||||||||||||||| |||||
Db       1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGTGCTCCTGGGCATAGTCCTCGCCGCCTGGCTT 60


Qy     106 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 165
           || || |||||||||||||||||||||||| ||||||||| ||||||||| |||||||||||
Db      61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120


Qy     166 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 225
           |||||||| ||||||||||||||| |||| |||||| ||||||| |||||||||||||||||
Db     121 GACCTGCTGCCCCACTTCCTGGTAGAGCCTGAGGACGTGTACATTGTCAAGAACAAGCCG 180


Qy     226 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 285
           ||| || | |||||||||||| ||||| ||||| ||||||||||||||||||||| |||||
Db     181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240


Qy     286 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 345
           ||||| ||||||||| ||||| || || || ||||||||| ||| | ||||| || |||||
Db     241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300


Qy     346 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 405
           |||||||||||||| || || || || ||||||||||| ||||| |||||| |||||||||
Db     301 ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG 360


Qy     406 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 465
           ||||||||||||||||| |||||||||||||||||||||| |||||||| ||||||||||||
Db     361 GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC 420
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Qy    466 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 525
          ||||||||| || |||    ||||||||||||| ||||||||||| |||||||||| |||
Db    421 TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG 480

Qy    526 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 585
          || ||||||||| ||||| || || || || || || || || || ||||| ||||| |||
Db    481 TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCCAGAAGGAATCCCCCCAGCTGAG 540

Qy    586 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 645
          |||||||||| || || ||||||| ||||||||| ||||| || ||||||||| ||||||
Db    541 GTGGAGTGGCTTCGAAATGAGGACCTCGTGGACCCCTCCCTCGATCCCAATGTGTACATC 600

Qy    646 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 705
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Db    601 ACGCGGGAGCACAGCCTAGTCGTGCGTCAGGCCCGCCTGGCCGACACGGCCAACTACACC 660

Qy    706 TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC 765
          || ||||||||||||||||||| || ||||||| ||| |||| || || ||||| || ||
Db    661 TGTGTGGCCAAGAACATCGTAGCCCGTCGCCGAAGCACCTCTGCAGCGGTCATTGTTTAT 720

Qy    766 GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC 825
          ||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||||
Db    721 GTGAACGGTGGGTGGTCGACGTGGACTGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGT 780

Qy    826 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC 885
          ||||||||||||||||||||||||||||||||||||||| |||||||||||||||| |||
Db    781 GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC 840

Qy    886 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC 945
          |||||||||||||||||||||||||||||||||||||| |||||||||||| || || |||
Db    841 TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACTCTGTGCCCAGTGGATGGGAGC 900

Qy    946 TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT 1005
          |||||  ||||||| ||||||| ||||||||| |||||||||||||||||||||||||||
Db    901 TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC 960

Qy   1006 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC 1065
          ||||||||||||||||||||||||||| |||| ||||||| | ||| ||||||||||||
Db    961 GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC 1020

Qy   1066 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC 1125
          |||||||||||||||||||||||||||| | |||| |||||| |||| ||||||||||
Db   1021 ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT 1080

Qy   1126 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC 1185
          ||||| | |||||| |||| .||||| || |||||| || || |||||| |||| | |||
Db   1081 CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA 1140

Qy   1186 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC 1245
          ||| |||| || || ||||||||| ||||||||| |||||||| ||||||||||||| |||
Db   1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC 1200

Qy   1246 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC 1305
          |||||  ||||||||||| ||||||||||||||||||||||||||||||||||| ||||||
Db   1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC 1260

Qy   1306 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG 1365
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                   |||||||||||  ||||||||||||||||||  ||||||||||||||||||||||  ||| |  ||
Db   1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG 1320

Qy   1366 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT 1425
                   ||||||||  |||||||||||||||||||||| | |||||| ||||||||||||||| |||||
Db   1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG 1380

Qy   1426 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC 1485
                   | |||||||| ||  ||||||||||||| |||||||||||||| ||||| |||||||||
Db   1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC 1440

Qy   1486 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT 1545
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Db   1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC 1500

Qy   1546 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC 1605
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Db   1501 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC 1560

Qy   1606 ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG 1665
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Db   1561 ATACCCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG 1620

Qy   1666 CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC 1725
                   || ||||||||||||||||||||||||||||||||||||||||||||||||  ||||||||
Db   1621 CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC 1680

Qy   1726 TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG 1785
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Db   1681 TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA 1740

Qy   1786 GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG 1845
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Db   1741 GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG 1800

Qy   1846 GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG 1905
                   ||||||||||||||||||| || ||||||| | || |||||||||||||||||||||||||||
Db   1801 GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG 1860

Qy   1906 GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC 1965
                   ||| | |||||||| ||||||||| |||||||||||||||||||||||||| |||||||||
Db   1861 GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC 1920

Qy   1966 CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC 2025
                   ||||||||||||||||| ||||||||||||| ||| |||||||||||| || ||||||| ||
Db   1921 CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG 1980

Qy   2026 TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG 2085
                   ||||| |||||||||||||| || ||||||| || ||||||||||| || |||||||||
Db   1981 TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG 2040

Qy   2086 GTGGTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC 2145
                   |||||||||||||||||||||||| || ||||||||||||||||||||| || |||||||||
Db   2041 GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC 2100

Qy   2146 TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG 2205
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Db      2101 TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG 2160

Qy      2206 AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG 2265
             ||||| ||||| ||||||||||||||||||||||||| ||||| |||||||||| |||||
Db      2161 AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCTTTTTACCACATCTGGAACGGCACC 2220

Qy      2266 CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG 2325
             |||| |||  |||||||||||||||||||||||||||  ||| | ||||||| || ||||||
Db      2221 CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG 2280

Qy      2326 GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC 2385
             |||||||||| ||||||||||||||||||||||||| || ||||||||||||| |||||||||
Db      2281 GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC 2340

Qy      2386 AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC 2445
             |||||||| ||||||||||||||||||||||  || ||||||||||||||||||| |||||||
Db      2341 AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy      2446 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2505
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||| ||||| ||
Db      2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy      2506 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2565
             ||||  |||||||||||||| || |||||  |||||||||| | ||||| ||||||||||||
Db      2461 GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy      2566 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2625
             |||||||||||||||||| ||||||||||||||||||||||||| ||||||||||||||||||||
Db      2521 CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy      2626 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2685
             ||||| ||||||||| |||||||||||||||||||||||||||| ||||||||| ||||| |||
Db      2581 AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy      2686 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGAGGC 2745
             || |||||||||||||||| ||||| |||||||||||||||||||||||| ||||| |||| |
Db      2641 GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGAGAC 2700

Qy      2746 CGGCCAG 2752
             | |||||
Db      2701 CAGCCAG 2707


RESULT 8
US-10-240-154-15
; Sequence 15,. Application US/10240154
; Publication No. US20030175741A1
; GENERAL INFORMATION:
;  APPLICANT: Cochran et al.
;  TITLE OF INVENTION: SCHIZOPHRENIA RELATED GENES
;  FILE REFERENCE: CKFW-P01-006
;  CURRENT APPLICATION NUMBER: US/10/240,154
;  CURRENT FILING DATE:  2001-04-02
;  PRIOR APPLICATION NUMBER: PCT/GB01/01486
;  PRIOR FILING DATE: 2001-04-02
;  NUMBER OF SEQ ID NOS: 34
;  SOFTWARE: PatentIn version 3.2
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    Query Match              81.8%;   Score 2252.2;   DB 16;   Length 2697;
    Best Local Similarity    89.7%;   Pred. No. 0;
    Matches 2419;  Conservative    0;  Mismatches  278;  Indels    0;  Gaps    0;

Qy        46 ATGGCCGTCCGGCCCGGCCTGTGGCCAGCGCTCCTGGGCATAGTCCTCGCCGCTTGGCTC 105
             |||||||||||||||||||||||||||| ||||||||||||||||||||||||| |||||
Db         1 ATGGCCGTCCGGCCCGGCCTGTGGCCAGTGCTCCTGGGCATAGTCCTCGCCGCCTGGCTT 60

Qy       106 CGCGGCTCGGGTGCCCAGCAGAGTGCCACCGTGGCCAACCCAGTGCCTGGTGCCAACCCG 165
             || || |||||||||||||||||||||||| ||||||||| ||||||||| |||||||||
Db        61 CGTGGTTCGGGTGCCCAGCAGAGTGCCACGGTGGCCAATCCAGTGCCCGGTGCCAACCCC 120

Qy       166 GACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATGTGTACATCGTCAAGAACAAGCCA 225
             |||||||| ||||||||||||||| |||||| |||| ||||||| |||||||||||||||
Db       121 GACCTGCTGCCCCACTTCCTGGTAGAGCCTGAGGACGTGTACATTGTCAAGAACAAGCCG 180

Qy       226 GTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGATCTTCTTCAAGTGCAACGGGGAG 285
             ||| || | |||||||||||| ||||| ||||| |||||||||||||||||||| |||||
Db       181 GTGTTGTTGGTGTGCAAGGCTGTGCCTGCCACCCAGATCTTCTTCAAGTGCAATGGGGAA 240

Qy       286 TGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCACAGACGGGAGCAGTGGGCTGCCC 345
             ||||| ||||||||| ||||| || || ||||||||| ||| | |||||| ||   ||||
Db       241 TGGGTCCGCCAGGTCGATCACGTAATTGAACGCAGCACCGACAGCAGCAGCGGATTGCCA 300

Qy       346 ACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGGTCGAGAAGGTGTTCGGGCTGGAG 405
             |||||||||||||| || || || || |||||||||| ||||| ||||| |||||||||
Db       301 ACCATGGAGGTCCGTATCAACGTATCGAGGCAGCAGGTAGAGAAAGTGTTTGGGCTGGAG 360

Qy       406 GAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGGGCACCACCAAGAGTCAGAAGGCC 465
             ||||||||||||||||| ||||||||||||||||||||| |||||||| |||||||||||
Db       361 GAATACTGGTGCCAGTGTGTGGCATGGAGCTCCTCGGGTACCACCAAAAGTCAGAAGGCC 420

Qy       466 TACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGCAGGAGCCGCTGGCCAAGGAGGTG 525
             |||||||| || |||     |||||||||||||||| ||||||||| ||||||||| |||
Db       421 TACATCCGGATTGCCTATTTGCGCAAGAACTTTGAGCAGGAGCCACTGGCCAAGGAAGTG 480

Qy       526 TCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCACCGGAGGGCATCCCTCCAGCCGAG 585
             || ||||||||| |||||| ||||| || || || || || |||| || || ||||| ||
Db       481 TCACTGGAGCAAGGCATTGTACTACCTTGTCGCCCCCCAGAAGGAATCCCCCCAGCTGAG 540

Qy       586 GTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGTCCCTGGACCCCAATGTATACATC 645
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Db       541 GTGGAGTGGCTTCGAAATGAGGACCTCGTGGACCCCTCCCTCGATCCCAATGTGTACATC 600

Qy       646 ACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCCTTGCTGACACGGCCAACTACACC 705
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Db       601 ACGCGGGAGCACAGCCTAGTCGTGCGTCAGGCCCGCCTGGCCGACACGGCCAACTACACC 660
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Qy    706  TGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCGCCTCCGCTGCTGTCATCGTCTAC  765
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Db    661  TGTGTGGCCAAGAACATCGTAGCCCGTCGCCGAAGCACCTCTGCAGCGGTCATTGTTTAT  720

Qy    766  GTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGC  825
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Db    721  GTGAACGGTGGGTGGTCGACGTGGACTGAGTGGTCCGTCTGCAGCGCCAGCTGTGGGCGT  780

Qy    826  GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCGCCTCTCAACGGGGGCGCTTTC  885
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Db    781  GGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGGCACCTCTCAACGGGGGCGCCTTC  840

Qy    886  TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACCCTGTGCCCAGTAGACGGCAGC  945
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Db    841  TGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCACTCTGTGCCCAGTGGATGGGAGC  900

Qy    946  TGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGGACTGCACCCACTGGCGGAGCCGT  1005
           |||||    |||||||| |||||||| |||||||| |||||||||||||||||||||||||
Db    901  TGGAGTTCGTGGAGTAAGTGGTCAGCCTGTGGGCTTGACTGCACCCACTGGCGGAGCCGC  960

Qy    1006 GAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGGAGTGCCAGGGCACTGACCTGGAC  1065
           ||||||||||||||||||||||||||| ||||| |||||||| | |||   |||||||||||
Db    961  GAGTGCTCTGACCCAGCACCCCGCAATGGAGGTGAGGAGTGTCGGGGTGCTGACCTGGAC  1020

Qy    1066 ACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCC  1125
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Db    1021 ACCCGCAACTGTACCAGTGACCTCTGCCTGCACACCGCTTCTTGCCCCGAGGACGTGGCT  1080

Qy    1126 CTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATC  1185
           ||||| | |||||| |||| |||| |||||| || |||||  || |||||  | |||
Db    1081 CTCTACATCGGCCTTGTCGCTGTGGCTGTGTGCCTCTTCTTGCTGTTGCTGGCCCTTGGA  1140

Qy    1186 CTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTC  1245
           ||| |||| || || |||||||| ||||||||| ||||||||| ||||||||||| |||
Db    1141 CTCATTTACTGTCGCAAGAAGGAAGGGCTGGACTCCGATGTGGCCGACTCGTCCATCCTC  1200

Qy    1246 ACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCATCTGCTC  1305
           ||||| |||||||||||| ||||||||||||||||||||||||||||||||||| ||||||
Db    1201 ACCTCGGGCTTCCAGCCTGTCAGCATCAAGCCCAGCAAAGCAGACAACCCCCACCTGCTC  1260

Qy    1306 ACCATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGG  1365
           ||||||||||| ||||||||||||||||||| |||||||||||||||||| ||  |   ||
Db    1261 ACCATCCAGCCAGACCTCAGCACCACCACTACCACCTACCAGGGCAGTCTATGTTCGAGG  1320

Qy    1366 CAGGATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGT  1425
           |||||||| ||||||||||||||||||||| |  |||| | |||||||||||||| |||||
Db    1321 CAGGATGGACCCAGCCCCAAGTTCCAGCTCTCTAATGGTCACCTGCTCAGCCCACTGGGG  1380

Qy    1426 GGCGGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCC  1485
           | ||||||||| || ||||||||||||||| ||||||||||||| ||||| |||||||||
Db    1381 AGTGGCCGCCATACGTTGCACCACAGCTCACCCACCTCTGAGGCTGAGGACTTCGTCTCC  1440

Qy    1486 CGCCTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTAT  1545
           ||||||||||||||| |||||||| || |||||||||||||| ||||||||||||| ||||
Db    1441 CGCCTCTCCACCCAAAACTACTTTCGTTCCCTGCCCCGCGGCACCAGCAACATGGCCTAC  1500

Qy    1546 GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTC  1605
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Db     1501  GGGACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACGGGGATCAGCCTCCTC  1560

Qy     1606  ATCCCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAG  1665
              ||  |||||| |||||||||| |||||||| |||||||||  ||||||||||||||||  |||||||||
Db     1561  ATACCCCCGGATGCCATCCCCCGAGGAAAGATCTACGAGATCTACCTCACACTGCACAAG  1620

Qy     1666  CCGGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGC  1725
              ||  |||||||||||||||||||||||||||||||||||||||||||||||||||||||    ||||||||
Db     1621  CCAGAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCAGTCGTTAGC  1680

Qy     1726  TGTGGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGG  1785
              |||||  ||||||  || |||||||||||||||||||||||||||| ||  ||||||||||||||
Db     1681  TGTGGGCCCCCAGGAGTCCTGCTCACCCGGCCAGTCATCCTTGCAATGGACCACTGTGGA  1740

Qy     1786  GAGCCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGG  1845
              |||||||||||||||||||||||||||| ||||||||||||||||||| |||||||||||| |||
Db     1741  GAGCCCAGCCCTGACAGCTGGAGTCTGCGCCTCAAAAAGCAGTCCTGCGAGGGCAGTTGG  1800

Qy     1846  GAGGATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAG  1905
              |||||||||||||||||||| ||  ||||||| |  ||  ||||||||||||||||||||||||||
Db     1801  GAGGATGTGCTGCACCTTGGTGAGGAGTCACCTTCCCACCTCTACTACTGCCAGCTGGAG  1860

Qy     1906  GCCAGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCC  1965
              |||  | |||||||||| |||||||||| |||||||||||||||||||||||||||| |||||||||
Db     1861  GCCGGGGCCTGCTATGTCTTCACGGAGCAGCTGGGCCGCTTTGCCCTGGTAGGAGAGGCC  1920

Qy     1966  CTCAGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACC  2025
              |||||||||||||||||  |||||||||||| ||| |||||||||| || ||||||||  ||
Db     1921  CTCAGCGTGGCTGCCACCAAGCGCCTCAGGCTCCTTCTGTTTGCTCCCGTGGCCTGTACG  1980

Qy     2026  TCCCTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAG  2085
              |||||  ||||||||||||||| || |||||||| ||  |||||||||||||  |||||||||
Db     1981  TCCCTTGAGTACAACATCCGAGTGTACTGCCTACACGACACCCACGACGCTCTCAAGGAG  2040

Qy     2086  GTGGTGCAGCTGGAGAAGCAGCTGGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCAC  2145
              ||||||||||||||||||||||||||  || ||||||||||||||||||||  |||||||||
Db     2041  GTGGTGCAGCTGGAGAAGCAGCTAGGTGGACAGCTGATCCAGGAGCCTCGCGTCCTGCAC  2100

Qy     2146  TTCAAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGG  2205
              |||||  |||||||||||||||||  || ||  |||||||||| |||||||||||||||||
Db     2101  TTCAAAGACAGTTACCACAACCTACGTCTCTCCATCCACGACGTGCCCAGCTCCCTGTGG  2160

Qy     2206  AAGAGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACG  2265
              |||||  |||||  |||||||||||||||||||||||| ||||| |||||||||||||  |||||
Db     2161  AAGAGCAAGCTACTTGTCAGCTACCAGGAGATCCCCTTTTTACCACATCTGGAACGGCACC  2220

Qy     2266  CAGCGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTG  2325
              ||||  |||   |||||||||||||||||||||||||  |||  | |||||||| || |||||||
Db     2221  CAGCAGTATCTGCACTGCACCTTCACCCTGGAGCGCATCAACGCCAGCACCAGCGACCTG  2280

Qy     2326  GCCTGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTC  2385
              |||||||||  |||||||||||||||||||||||||  || ||||||||||||||||| |||||||||||
Db     2281  GCCTGCAAGGTGTGGGTGTGGCAGGTGGAGGGAGATGGGCAGAGCTTCAACATCAACTTC  2340

Qy     2386  AACATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTC  2445
              |||||||||  ||||||||||||||||||||||||||||||     ||  |||||||||||||||||||| |||||||
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Db     2341 AACATCACTAAGGACACAAGGTTTGCTGAATTGTTGGCTCTGGAGAGTGAAGGGGGGGTC 2400

Qy     2446 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATT 2505
            ||||||||||||||||||||||||||||||||||||||||||||||||||||| |||| ||
Db     2401 CCAGCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAAAAGATCATC 2460

Qy     2506 TCCAGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTC 2565
            ||||  |||||||||||||| || ||||| ||||||||| | |||||  |||||||||||
Db     2461 GCCAGTCTGGACCCACCCTGCAGCCGGGGCGCCGACTGGAGAACTCTAGCCCAGAAACTT 2520

Qy     2566 CACCTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTC 2625
            ||||||||||||||||| ||||||||||||||||||||||||| |||||||||||||||||
Db     2521 CACCTGGACAGCCATCTTAGCTTCTTTGCCTCCAAGCCCAGCCCTACAGCCATGATCCTC 2580

Qy     2626 AACCTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTG 2685
            |||||  ||||||||  ||||||||||||||||||||||  ||||||||| ||||| |||
Db     2581 AACCTATGGGAGGCACGGCACTTCCCCAACGGCAACCTCGGCCAGCTGGCAGCAGCTGTG 2640

Qy     2686 GCTGGACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
            || |||||||||| ||||| |||||||||||||||| |||||||||| ||||| |||
Db     2641 GCCGGACTGGGCCAACCAGATGCTGGCCTCTTCACGGTGTCGGAGGCCGAGTGTTGA 2697
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RESULT 9
US-09-933-261-2
; Sequence 2, Application US/09933261
; Publication No. US20030040046A1
;   GENERAL INFORMATION:
;   APPLICANT: Tessier-Lavigne, Marc
;   Leonardo, E. David
;   Hink, Lindsay
;   Masu, Masayuki
;   Kazuko, Keino-Masu
;   TITLE OF INVENTION: Netrin Receptors
;   NUMBER OF SEQUENCES: 8
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;   STREET: 268 BUSH STREET, SUITE 3200
;   CITY: SAN FRANCISCO
;   STATE: CALIFORNIA
;   COUNTRY: USA
;   ZIP: 94104
;   COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/09/933,261
;   FILING DATE: 20-Aug-2001
;   CLASSIFICATION: <Unknown>
;   PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: 08/808,982
;   FILING DATE: <Unknown>
;   ATTORNEY/AGENT INFORMATION:
;   NAME: OSMAN, RICHARD A
;   REGISTRATION NUMBER: 36,627

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   Query Match              56.8%;  Score 1562.4;  DB 10;  Length 1787;
   Best Local Similarity    98.5%;  Pred. No. 0;
   Matches 1661;  Conservative    0;  Mismatches   16;  Indels    9;  Gaps    8;


Qy      1070 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1129
             |||||||||||||||||||||||| ||||||| |||||||||||||||||||||||||||
Db         1 GCAACTGTACCAGTGACCTCTG-GTACACACTGCTTCTGGCCCTGAGGACGTGGCCCTCT 59


Qy      1130 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 1189
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119


Qy      1190 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1249
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       120 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 179


Qy      1250 CAGGCTTCCAGCCCGTCAGCATC-AAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 1308
             ||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Db       180 CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239


Qy      1309 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1368
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       240 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299


Qy      1369 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1428
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       300 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359


Qy      1429 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1488
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       360 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419


Qy      1489 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 1548
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       420 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 479


Qy      1549 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1608
             ||||||||||||||||||||||||||||||||||||||||||||||| |||||||||||||
Db       480 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATC 539


Qy      1609 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1668
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       540 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599
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Qy    1669 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1728
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     600 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy    1729 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1788
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     660 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy    1789 CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1848
           ||||||||||||||||||||||||||    ||||||||||||||||||||||| |||||||||
Db     720 CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy    1849 GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1908
           |||||  |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     779 GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy    1909 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1968
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     838 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy    1969 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 2028
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     898 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957

Qy    2029 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2088
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     958 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017

Qy    2089 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTC 2148
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    1018 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

Qy    2149 AAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAG 2208
           ||||||||||||||||||   |||         ||||||||||||||||||||||||||||
Db    1077 AAGGACAGTTACCACAACCT--GCCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy    2209 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2268
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1135 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy    2269 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2328
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1195 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy    2329 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2388
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1255 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 1314

Qy    2389 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 2448
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1315 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 1374

Qy    2449 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 2508
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1375 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 1434
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Qy    2509 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2568
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1435 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 1494

Qy    2569 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 2628
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1495 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

Qy    2629 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 2688
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1555 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 1614

Qy    2689 GGACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGAGGCC 2746
           ||      |||||||  ||||  ||||  ||  |||||||||  ||||||||||||||||||
Db    1615 GGGACTGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGAGGCC 1674

Qy    2747 GGCCAG 2752
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Db    1675 GGCCAG 1680


RESULT 10
US-10-256-702-2
; Sequence 2, Application US/10256702
; Publication No. US20030059859A1
;    GENERAL INFORMATION:
;         APPLICANT: Tessier-Lavigne, Marc
;                    Leonardo, E. David
;                    Hink, Lindsay
;                    Masu, Masayuki
;                    Kazuko, Keino-Masu
;         TITLE OF INVENTION: Netrin Receptors
;         NUMBER OF SEQUENCES: 8
;         CORRESPONDENCE ADDRESS:
;              ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
;              STREET: 268 BUSH STREET, SUITE 3200
;              CITY: SAN FRANCISCO
;              STATE: CALIFORNIA
;              COUNTRY: USA
;              ZIP: 94104
;         COMPUTER READABLE FORM:
;              MEDIUM TYPE: Floppy disk
;              COMPUTER: IBM PC compatible
;              OPERATING SYSTEM: PC-DOS/MS-DOS
;              SOFTWARE: PatentIn Release #1.0, Version #1.30
;         CURRENT APPLICATION DATA:
;              APPLICATION NUMBER: US/10/256,702
;              FILING DATE: 27-Sep-2002
;              CLASSIFICATION: <Unknown>
;         PRIOR APPLICATION DATA:
;              APPLICATION NUMBER: US/09/933,261
;              FILING DATE: 20-Aug-2001
;              APPLICATION NUMBER: 08/808,982
;              FILING DATE: <Unknown>
;         ATTORNEY/AGENT INFORMATION:
;              NAME: OSMAN, RICHARD A
;              REGISTRATION NUMBER: 36,627
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  Query Match              56.8%;   Score 1562.4;   DB 14;   Length 1787;
  Best Local Similarity    98.5%;   Pred. No. 0;
  Matches 1661;  Conservative     0;  Mismatches   16;  Indels    9;  Gaps     8;


Qy      1070 GCAACTGTACCAGTGACCTCTGTGTACACAGTGCTTCTGGCCCTGAGGACGTGGCCCTCT 1129
             |||||||||||||||||||||||||| |||||||| |||||||||||||||||||||||||
Db         1 GCAACTGTACCAGTGACCTCTG-GTACACACTGCTTCTGGCCCTGAGGACGTGGCCCTCT 59


Qy      1130 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 1189
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        60 ATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGGTCCTGCTGCTGCTTGTCCTCATCCTCG 119


Qy      1190 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 1249
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       120 TTTATTGCCGGAAGAAGGAGGGGCTGGACTCAGATGTGGCTGACTCGTCCATTCTCACCT 179


Qy      1250 CAGGCTTCCAGCCCGTCAGCATC-AAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 1308
             |||||||||||||||||||||||| |||||||||||||||||||||||||||||||||||
Db       180 CAGGCTTCCAGCCCGTCAGCATCTAAGCCCAGCAAAGCAGACAACCCCCATCTGCTCACC 239


Qy      1309 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 1368
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       240 ATCCAGCCGGACCTCAGCACCACCACCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAG 299


Qy      1369 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 1428
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       300 GATGGGCCCAGCCCCAAGTTCCAGCTCACCAATGGGCACCTGCTCAGCCCCCTGGGTGGC 359


Qy      1429 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 1488
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       360 GGCCGCCACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGC 419


Qy      1489 CTCTCCACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGG 1548
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy      1549 ACCTTCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATC 1608
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Qy      1609 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 1668
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Db       540 CCCCCAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCG 599
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Qy    1669 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 1728
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db     600 GAAGACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGT 659

Qy    1729 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 1788
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db     660 GGACCCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAG 719

Qy    1789 CCCAGCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAG 1848
           IIIIIIIIIIIIIIIIIIIIIIIIIIII IIIIIIIIIIIIIIIIIIIIIIIII IIIIIIIII
Db     720 CCCAGCCCTGACAGCTGGAGCCTGGCCCTCAAAAAGCAGTCGTGCGAGGG-AGCTGGGAG 778

Qy    1849 GATGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 1908
           IIIII IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db     779 GATGT-CTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCC 837

Qy    1909 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 1968
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db     838 AGTGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTC 897

Qy    1969 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 2028
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db     898 AGCGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCC 957

Qy    2029 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 2088
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db     958 CTCGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTG 1017

Qy    2089 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTC 2148
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db    1018 GTGCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTT- 1076

Qy    2149 AAGGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAG 2208
           IIIIIIIIIIIIIIIIIIIIII III IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db    1077 AAGGACAGTTACCACAACCT--GCCTATCATCCACGATGTGCCCAGCTCCCTGTGGAAG 1134

Qy    2209 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 2268
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db    1135 AGTAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAG 1194

Qy    2269 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 2328
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db    1195 CGGTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCC 1254

Qy    2329 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 2388
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db    1255 TGCAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAAC 1314

Qy    2389 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 2448
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db    1315 ATCACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCA 1374

Qy    2449 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 2508
           IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
Db    1375 GCCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCC 1434
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Qy      2509 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 2568
             ||||||||||||||||||||||||||||||||||||||||||||||.||||||||||||||
Db      1435 AGCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCAC 1494

Qy      2569 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 2628
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1495 CTGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAAC 1554

Qy      2629 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 2688
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1555 CTGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT 1614

Qy      2689 GGACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGAGGCC 2746
             ||     |||||||   ||||  ||||  || ||||||||| |||||||||||||||||||
Db      1615 GGGACTGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGAGGCC 1674

Qy      2747 GGCCAG 2752
             ||||||
Db      1675 GGCCAG 1680


RESULT 11
US-10-296-115-365
; Sequence 365, Application US/10296115
; Publication No. US20040053248A1
; GENERAL INFORMATION:
;  APPLICANT: Hyseq Inc
;  TITLE OF INVENTION: No. US20040053248A1el Nucleic Acids and Polypeptides
;  FILE REFERENCE: 784PCT
;  CURRENT APPLICATION NUMBER: US/10/296,115
;  CURRENT FILING DATE:  2002-11-18
;  PRIOR APPLICATION NUMBER: US09/488,725
;  PRIOR FILING DATE: 2000-01-21
;  PRIOR APPLICATION NUMBER: US09/552,317
;  PRIOR FILING DATE: 2000-04-25
;  NUMBER OF SEQ ID NOS: 1478
; SEQ ID NO 365
;   LENGTH: 1321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-296-115-365

  Query Match           43.8%;  Score 1206.6;  DB 17;  Length 1321;
  Best Local Similarity  98.0%;  Pred. No. 1.2e-299;
  Matches 1295;  Conservative    0;  Mismatches   19;  Indels    7;  Gaps     7;

Qy      1435 CACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCC 1494
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         1 CACACACTGCACCACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCC 60

Qy      1495 ACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTC 1554
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        61 ACCCAGAACTACTTCCGCTCCCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCTTC 120

Qy      1555 AACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCCCA 1614
             |||||||||||||||||||||||||||||||||||||||| |||||||||||||||||||
Db       121 AACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGAATCAGCCTCCTCATCCCCCCA 180
```

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Qy    1615 GATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGAC 1674
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     181 GATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAGAC 240

Qy    1675 GTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCC 1734
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     241 GTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGACCC 300

Qy    1735 CCT-GGCGTCCTGCTCACCCGGCCAGTCATCCT-GGCTATGGACCACTGT-GGGGAGCCC 1791
           ||| ||||||||||| ||||||||||||||||| || ||||||||||||| |||||||||
Db     301 CCTGGGCGTCCTGCTTACCCGGCCAGTCATCCTGGGTATGGACCACTGTGGGGGAGCCC 360

Qy    1792 AGCCCTGACAGCT-GGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGA 1850
           ||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db     361 AGCCCTGACAGCTGGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGA 420

Qy    1851 TGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAG 1910
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     421 TGTGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAG 480

Qy    1911 TGCCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAG 1970
           |||||||||||||||||||||||||||||| |||||| ||||||||||||||||||||||
Db     481 TGCCTGCTACGTCTTCACCGAGCAGCTGAGCCGCTATGCCCTGGTGGGAGAGGCCCTCAG 540

Qy    1971 CGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCT 2030
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     541 CGTGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCT 600

Qy    2031 CGAGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGT 2090
           |||||||||||| | ||||||||||||||||||||| ||||||||||||||| |  |||||
Db     601 CGAGTACAACATACTGGTCTACTGCCTGCATGACACTCACGATGCACTCAACGTAGTGGT 660

Qy    2091 GCAGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAA 2150
           ||||||||||||||||||||    ||||||||||||||||||||||||| ||| ||||||||||
Db     661 GCAGCTGGAGAAGCAGCTGCAGGGACAGCTGATCCAGGAGCCACTGGTACTGCACTTCAA 720

Qy    2151 GGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAG 2210
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     721 GGACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAG 780

Qy    2211 TAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCG 2270
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     781 TAAGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCG 840

Qy    2271 GTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTG 2330
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     841 GTACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTG 900

Qy    2331 CAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACAT 2390
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     901 CAAGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACAT 960

Qy    2391 CACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGC 2450
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     961 CACCAAGGACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAGC 1020
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Qy      2451 CCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAG 2510
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1021 CCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCAG 1080

Qy      2511 CCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCT 2570
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1081 CCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACCT 1140

Qy      2571 GGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCT 2630
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1141 GGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACCT 1200

Qy      2631 GTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCT-G 2689
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db      1201 GTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTGG 1260

Qy      2690 GACTGGGCCAGCCAGACGCTGGCCTC-TTCACAGTG-TCGGAGGCTGAGTGCTGAGGCCG 2747
             ||||||||||||  ||||  ||||  ||  |||||||||| ||||||||||||||||||||
Db      1261 GACTGGGCCAGCAGGACGGTGGCTTCTTTCACAGTGTTCGGAGGCTGAGTGCTGAGGCCG 1320

Qy      2748 G 2748
             |
Db      1321 G 1321
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RESULT 12
US-10-087-684-1
; Sequence 1, Application US/10087684
; Publication No. US20040029116A1
; GENERAL INFORMATION:
;   APPLICANT: Edinger, Shlomit R.
;   APPLICANT: MacDougall, John R.
;   APPLICANT: Millet, Isabelle
;   APPLICANT: Ellerman, Karen
;   APPLICANT: Stone, David J.
;   APPLICANT: Grosse, William M.
;   APPLICANT: Lepley, Denise M.
;   APPLICANT: Rieger, Daniel K.
;   APPLICANT: Burgess, Cathereine E.
;   APPLICANT: Casman, Stacie, J.
;   APPLICANT: Spytek, Kimberly A.
;   APPLICANT: Boldog, Ferenc L.
;   APPLICANT: Li, Li
;   APPLICANT: Padigaru, Muralidhara
;   APPLICANT: Mishra, Vishnu
;   APPLICANT: Shenoy, Suresh G. ...
;   APPLICANT: Rastelli, Luca
;   APPLICANT: Tchernev, Velizar T.
;   APPLICANT: Vernet, Corine A.M.
;   APPLICANT: Zerhusen, Bryan D.
;   APPLICANT: Malyankar, Uriel M.
;   APPLICANT: Guo, Xiaojia
;   APPLICANT: Miller, Charles E.
;   APPLICANT: Gangolli, Esha A.
;   TITLE OF INVENTION: PROTEINS AND NUCLEIC ACIDS ENCODING SAME
;   FILE REFERENCE: 21402-214 CIP
;   CURRENT APPLICATION NUMBER: US/10/087,684

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  Query Match                34.0%;   Score 936.2;  DB 17;  Length 2860;
  Best Local Similarity      61.7%;   Pred. No. 3.6e-230;
  Matches 1662;  Conservative     0;  Mismatches  938;  Indels   93;  Gaps     7;

Qy        143 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 202
              || | | || |     ||     | || | ||| ||| |||||||| |||||| |||| |
Db        168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227

Qy        203 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 262
                |||||| || |||||||||||| ||| ||| ||| ||| | ||||||||| ||||
Db        228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTCCGCTGCCGCGCCTTCCCCGCCACACAGA 287

Qy        263 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 322
              ||| |||||||||||||||||||| |||||||| ||||| ||||||| | || ||
Db        288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347

Qy        323 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 382
              || | | || || |||| |||| | || | || || ||||||||||
Db        348 TGGATGAGGCCACCGGTCTGCGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407

Qy        383 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 442
              | ||| || | || |||||||||||| ||||||||||||||||||||.|| ||||||||| |||
Db        408 TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCGG 467

Qy        443 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 502
              |||||||||||||    ||||||| ||||||| |||     ||||||||||||||||| |
Db        468 GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527

Qy        503 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 562
              |||||||  ||||  |||||||||| ||||||| ||| || |   |  | |||| ||||| || |
Db        528 AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587

Qy        563 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 622
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                       |||||||  |  |||   |||||||||||| ||||||| ||| |||||| |  | |||||
Db    588 CGGAGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA 647

Qy    623 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 682
                ||| |||| |||| |      |||| || ||||| |||  |  | || |||||||||||
Db    648 CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC 707

Qy    683 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 742
                |  | |||||| ||||||||| ||||||||||||||||||||||||||||   ||||| |||
Db    708 TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA 767

Qy    743 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 802
                || | || | ||||||||||||||||||| || || |||||| |    ||| | ||||||||
Db    768 CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC 827

Qy    803 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 862
                |||| | | |||||||| || |||||||||||| || | |||||| ||||||||||||| |
Db    828 CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG 887

Qy    863 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 922
                | || |||||||| || || ||||| ||||| ||| |      ||||||| || ||||||| |||
Db    888 CTCCACTCAACGGAGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA 947

Qy    923 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 982
                || | ||||||||| || ||    |||| ||||||||||||||||||| |||||  |   |
Db    948 CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG 1007

Qy    983 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 1042
                | || ||||||||||| |||||| ||||| |||||| |  | || |||||| |||||||| |
Db   1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG 1067

Qy   1043 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1102
                | ||| || ||  || ||| | |||| |     ||||| ||  || || | ||  |   |
Db   1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG 1127

Qy   1103 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG 1162
                   |      || | | |        |  | |  | | | ||| | |    || | ||||
Db   1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG 1187

Qy   1163 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG 1222
                 | || ||   ||     |  | | || || |||| ||      |    | ||| |||
Db   1188 CAATCCTCATGGCGGTGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG 1247

Qy   1223 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA 1279
                | | |  |||||||| || | || || || || |||||| |||||| | | |||| |
Db   1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG 1307...

Qy   1280 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA 1333
                  |  |    |||||| || || || |  |  | || |||||| |   ||| |
Db   1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG 1367

Qy   1334 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA--------------------- 1370
                || || |||||| ||       | | | ||| ||||||
Db   1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA 1427

Qy   1371 ----------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1399
                                      ||||||| ||| || || || ||  ||
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Db   1428 CCAACTCTCCTCTGCTGGACCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA 1487

Qy   1400 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC---------ACTGCACC 1447
          | ||||| |   |  ||| ||    |  | ||| | |   |              | ||
Db   1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy   1448 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1507
          || | ||  | |  |  |  ||| ||| |   | || ||  |          |     | |
Db   1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy   1508 TCCGCTC---------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1552
          || | ||               ||||||||||| | |  | | ||    |  || ||||
Db   1608 TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT 1667

Qy   1553 TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC 1612
          |  || ||| || ||  |||| |   | ||||     ||||| |||| | || | | |||
Db   1668 TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA 1727

Qy   1613 CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG 1672
          | ||||| |||| | |  ||| ||| |||| | | |||| | ||  |  ||||| | |||
Db   1728 ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA 1787

Qy   1673 ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC 1732
           |  ||   || ||   ||  |||||     | ||||  |  ||  |    ||||||||||
Db   1788 GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC 1847

Qy   1733 CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA 1792
          || | |||  | ||||||  |   ||| || || | |||| | ||||||||| ||   ||
Db   1848 CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA 1907

Qy   1793 GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG 1852
          |  |     ||||| | | |   ||||| | |  ||| | | | |||||| ||||||||| |
Db   1908 GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG 1967

Qy   1853 TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG 1912
          || ||   ||||| |||||| | |     |    | ||||||||||||||||| ||||| |
Db   1968 TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG 2027

Qy   1913 CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG 1972
          ||||  || || |    ||  ||||||||||  || |   |   ||| ||| || |    |
Db   2028 CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC 2087

Qy   1973 TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG 2032
           | || | |||||| |||||||| ||| |||| |  | |  | | | ||||||||||| |
Db   2088 GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG 2147

Qy   2033 AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC 2092
          |||||| | ||||||||||||||||||||| | |||| | | |    ||||| ||||||||| ||
Db   2148 AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG 2207

Qy   2093 AGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG 2152
          |||||||| |   ||||| ||| |  || | ||||||||       || |||||||
Db   2208 AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG 2267

Qy   2153 ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA 2212
          |||||||||||||||||||||||| ||| |||| || | ||| |||| ||| |
Db   2268 ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA 2327
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```
Qy   2213 AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT 2272
          ||||  || | || |||||||||||||||| |||||||| |||| ||||| ||| |
Db   2328 AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG 2387

Qy   2273 ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA 2332
           |  | |||||||| |||||||||||| |    ||||    | | | || || ||||||
Db   2388 CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA 2447

Qy   2333 AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA 2392
          || | || ||| |||| ||||| || || || |||| |||   |  |  | ||  |
Db   2448 AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG 2507

Qy   2393 CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG 2449
          |  ||       | | |   ||   || ||   |  |     ||         ||| |
Db   2508 CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA 2567

Qy   2450 CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA 2509
          ||| | |||| ||  |||||||||||||||||||| |   ||| || |||||||||   ||
Db   2568 CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA 2627

Qy   2510 GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC 2569
          |||| || | |||     ||||| |||||||||| | || ||||| |||
Db   2628 GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA 2687

Qy   2570 TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC 2629
          ||||| |   | || |   | ||||||| |||| | |||||||| | |||||||  |||
Db   2688 TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC 2747

Qy   2630 TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG 2689
          | ||||| || | ||| | | || | || |||||| | |||||   |||| ||||  |
Db   2748 TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG 2807

Qy   2690 GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
          ||||| || || |   | | || | || ||||||
Db   2808 AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA 2860
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RESULT 13
US-10-087-684-3
; Sequence 3, Application US/10087684
; Publication No. US20040029116A1
; GENERAL INFORMATION:
; APPLICANT: Edinger, Shlomit R.
; APPLICANT: MacDougall, John R.
; APPLICANT: Millet, Isabelle
; APPLICANT: Ellerman, Karen
; APPLICANT: Stone, David J.
; APPLICANT: Grosse, William M.
; APPLICANT: Lepley, Denise M.
; APPLICANT: Rieger, Daniel K.
; APPLICANT: Burgess, Cathereine E.
; APPLICANT: Casman, Stacie, J.
; APPLICANT: Spytek, Kimberly A.
; APPLICANT: Boldog, Ferenc L.
; APPLICANT: Li, Li
; APPLICANT: Padigaru, Muralidhara

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   Query Match              34.0%;  Score 936.2;  DB 17;  Length 2860;
   Best Local Similarity    61.7%;  Pred. No. 3.6e-230;
   Matches 1662;  Conservative    0;  Mismatches  938;  Indels   93;  Gaps    7;

Qy        143 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 202
              || |  | || |    ||        | || | ||| ||| |||||||||  ||||||  |||| |
Db        168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227

Qy        203 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 262
              |||||  || ||||||||||||| ||||  |||| |   ||| ||||  | ||||||||| ||||
Db        228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTTCGCTGCCGCGCCTTCCCCGCCACACAGA 287

Qy        263 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 322
              |||  ||||||||||||||||||||| |||||||||  |||||    ||||||||| |   ||    ||
Db        288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347

Qy        323 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 382
              || | |  ||  ||  ||| |||||         ||||| |  || | || || |||||||||
Db        348 TGGATGAGGCCACCGGCCTGCGGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407
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Qy    383  TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 442
           |  |||  || |  ||  ||||||||||| ||||||||||||||||||||| |||||||||  | |
Db    408  TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCAG 467

Qy    443  GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 502
           |||||||||||||||   ||||||  |||||| |||   |||||||||||||||||| |
Db    468  GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527

Qy    503  AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 562
           ||||||| ||||  ||||||||||  |||||||||| |||||| || |     |  |  |||| ||||| || |
Db    528  AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587

Qy    563  CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 622
           |||||||   | |||     |||||||||| ||||||  |||  ||||||   |  | |||||
Db    588  CGGAGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA 647

Qy    623  CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 682
           ||| |||| ||||   |    |||| || |||| |||   |  | || |||||||||||
Db    648  CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC 707

Qy    683  TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 742
           |  |  |||||| |||||||| ||||||||||||||||||||||||||||||||     ||||| |||
Db    708  TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA 767

Qy    743  CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 802
           || | ||  | ||||||||||||||||| || || ||||| |   ||| | |||||||||
Db    768  CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC 827

Qy    803  TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 862
           ||||   |   | |||||||| ||  |||||||||||| || |  ||||| |||||||||||| |
Db    828  CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG 887

Qy    863  CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 922
           | || |||||||| ||  || ||  ||||| ||||| |||  |       ||||||| || ||||||| |||
Db    888  CTCCACTCAACGGAGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA 947

Qy    923  CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 982
           || | ||||||||| || ||     |||| ||||||||||||||||||| ||||| |     |
Db    948  CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG 1007

Qy    983  ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 1042
           | || ||||||||||| ||||| |||||| |  ||  ||||||  ||||||| |
Db    1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG 1067

Qy    1043 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1102
           | ||| || || |  .....|| ||| | ||||| || || || || || | || |
Db    1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG 1127

Qy    1103 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG 1162
           |          || | | ||          | | || | ||| |   ||| | |||
Db    1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG 1187

Qy    1163 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG 1222
           | || || || |   | | || || |||| || |   |  || |||
Db    1188 CAATCCTCATGGCGGTGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG 1247

Qy    1223 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA 1279
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           |  |   ||||||| ||    |    || ||     || |||||| ||||||| | | ||| |
Db   1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG 1307

Qy   1280 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA 1333
           |  |    ||||| || || ||    |  | | |  || |||||  |   ||| |
Db   1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG 1367

Qy   1334 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA----------------------- 1370
           ||  ||  ||||| ||     |  | | ||| ||||||
Db   1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA 1427

Qy   1371 ----------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1399
                                  ||||||| ||| || || ||   ||
Db   1428 CCAACTCTCCTCTGCTGGACCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA 1487

Qy   1400 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC----------ACTGCACC 1447
             | |||| |  |   ||| ||     |  | |||| |   |          |  ||
Db   1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy   1448 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1507
            ||  |  || | | |   |     ||| |||   |  || ||  |       |    | |
Db   1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy   1508 TCCGCTC---------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1552
            || | ||               ||||||||||| | |  | | | ||    | || ||||
Db   1608 TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT 1667

Qy   1553 TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC 1612
            |   || ||| || || || |||| | |||||     ||||| |||||| | || | |||
Db   1668 TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA 1727

Qy   1613 CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG 1672
            |  ||||| |||| || ||| |||| |||| |||||'|| || |      |||||| | |||
Db   1728 ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA 1787

Qy   1673 ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC 1732
              |  ||    ||  ||    |||||  | |||| | |||| ||| || | |||||||||
Db   1788 GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC 1847

Qy   1733 CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA 1792
            || | ||| |||||||   ||| || ||||||||| | ||| ||||||||| ||    ||
Db   1848 CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA 1907

Qy   1793 GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG 1852
            | |        ||||| |||||||  |||| | ||| |  |  | |||||| |||||||||| |
Db   1908 GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG 1967

Qy   1853 TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG 1912
            || ||  ||||| ||||||  |  |   |   | |  || ||||||||||||||||| |||| |
Db   1968 TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG 2027

Qy   1913 CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG 1972
            |||| || || | || |||||||||| ||  |   | | |   ||| ||| || |   |
Db   2028 CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC 2087

Qy   1973 TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG 2032
            | || | ||||||| ||| ||| |||||     | || || || |     |||||||||||| |
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Db    2088  GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG  2147

Qy    2033  AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC  2092
            ||||||  |  ||||||||||||||||||  |  ||||||  |  |   |||||  ||||||||||  ||
Db    2148  AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG  2207

Qy    2093  AGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG  2152
            ||||||||  |    |||||  |||  |  |  ||  |  |||||||        ||      |||||||
Db    2208  AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG  2267

Qy    2153  ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA  2212
            |||||||||||||||||||||||  |||  ||||  ||  |  |||      |||   ||||  |||  |
Db    2268  ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA  2327

Qy    2213  AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT  2272
            ||||  ||  |  ||   |||||||||||||||||||  |||||||||  ||||  |||||   |||   |
Db    2328  AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG  2387

Qy    2273  ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA  2332
             |  |  ||||||||  ||||||||||||  |      ||||      |   |   ||  ||   ||||||
Db    2388  CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA  2447

Qy    2333  AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA  2392
            ||  |  ||  |||  ||||  ||||  |||||  ||  ||  ||  ||||   |   |    ||    |
Db    2448  AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG  2507

Qy    2393  CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG  2449
            |  ||         |  |  |    ||    ||       |    ||         |||  |
Db    2508  CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA  2567

Qy    2450  CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA  2509
            |||  |  |||| ||     ||||||||||||||||||  |   ||| || |||||||||      ||
Db    2568  CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA  2627

Qy    2510  GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC  2569
            ||||  ||   |  |||      |||||         |||||||||  |  ||  |||||  |||
Db    2628  GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA  2687

Qy    2570  TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC  2629
            |||||  |    |  ||  |   |  |||||||| ||||    |  |||||||| |     |||||||||  |||
Db    2688  TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC  2747

Qy    2630  TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG  2689
             | ||||| ||  |  |||      |  |  || ||||||  |  |||||        ||  |||  |
Db    2748  TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG  2807

Qy    2690  GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA  2742
            |||||  ||   ||        |        |   | ||  |  || |||||||
Db    2808  AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA  2860


RESULT 14
US-10-218-779-1
; Sequence 1, Application US/10218779
; Publication No. US20040029222A1
; GENERAL INFORMATION:
;  APPLICANT: Edinger, Shlomit
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    Query Match   34.0%; Score 936.2; DB 17; Length 2860;
    Best Local Similarity   61.7%; Pred. No. 3.6e-230;
    Matches 1662; Conservative   0; Mismatches 938; Indels   93; Gaps   7;

Qy   143 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 202
                || | | || || | || | ||| ||| ||||||||| |||||| |||| |
Db   168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227

Qy   203 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 262

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                  ||||| || |||||||||||| ||| |||       |||    ||| | ||||||||| ||||
Db     228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTCCGCTGCCGCGCCTTCCCCGCCACACAGA 287

Qy     263 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 322
               ||| ||||||||||||||||| ||||||||| |||||  ||||||||| |  ||  ||
Db     288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347

Qy     323 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 382
               || | | || || ||||    ||||| | || | || || |||||||||
Db     348 TGGATGAGGCCACCGGTCTGCGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407

Qy     383 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 442
            | |||  || |  || ||||||||||||| |||||||||||||||||||| ||||||||| |||
Db     408 TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCGG 467

Qy     443 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 502
            ||||||||||||||     |||||| |||||| |||       ||||||||||||||||| |
Db     468 GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527

Qy     503 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 562
            |||||||  ||||  ||||||||||| ||||| |||||| || |   |  | | ||||  ||||| || |
Db     528 AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587

Qy     563 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 622
            ||||||| | ||| |||||||||| |||||| ||| ||||| |  | |||||
Db     588 CGGAGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA 647

Qy     623 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 682
            ||| |||| |||| |     ||||   || | |||| |||   |  | || ||||||||||
Db     648 CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC 707

Qy     683 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 742
            | | ||||||  |||||||| ||||||||||||||||||||||||||||||||     ||||| |||
Db     708 TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA 767

Qy     743 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 802
            || | || | |||||||||||||||||||| || || ||||| |   ||| | ||||||||
Db     768 CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC 827

Qy     803 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 862
               ||||  |  | ||||||| || ||||||||||| || | |||||  |||||||||||| |
Db     828 CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG 887

Qy     863 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 922
            | || ||||||||  || || ||||| ||||| |||       ||||||| || |||||| |||
Db     888 CTCCACTCAACGGAGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA 947

Qy     923 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 982
            || | ||||||||| || || ||    |||| ||||||||||||||||| |||||  |   |
Db     948 CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG 1007

Qy     983 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 1042
            | || |||||||||| |||||| ||||| |||||| | || ||||||| ||||||||| |
Db    1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG 1067

Qy    1043 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1102
            | ||| || || ||  ||      || |  | |||   || |  |  |  ||
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Db   1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG 1127

Qy   1103 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG 1162
               |    || |  | |         | | || | | ||| |   |||   | |||
Db   1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG 1187

Qy   1163 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG 1222
               | ||   ||   ||        |  | || ||  |||| ||     |  | ||| |||
Db   1188 CAATCCTCATGGCGGTGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG 1247

Qy   1223 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA 1279
             | |  ||||||| ||  |    || ||     || |||||| ||||||| | | ||| |
Db   1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG 1307

Qy   1280 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA 1333
             | |   |||||| || || || ||      | | || | || |||||| |   ||| |
Db   1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG 1367

Qy   1334 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA----------------------- 1370
             ||   ||  |||||  ||    ||   | | | |||  ||||||
Db   1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA 1427

Qy   1371 --------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1399
                               |||||||  |||  || ||  ||     ||
Db   1428 CCAACTCTCCTCTGCTGGACCCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA 1487

Qy   1400 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC----------ACTGCACC 1447
             | |||| |  |  | ||| ||     |  | ||| | |  |          | ||
Db   1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy   1448 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1507
             ||  |  || | |  |  |   ||| |||   |  || || ||       |     | |
Db   1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy   1508 TCCGCTC--------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1552
             || | ||              ||||||||||| | |  | | ||   |  || ||||
Db   1608 TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT 1667

Qy   1553 TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC 1612
             |   || ||| || || |||| |    |||||  |||||   ||||| ||||| | | |||
Db   1668 TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA 1727

Qy   1613 CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG 1672
             | ||||| |||| ||  | |||  |||| ||||| || ||  |   |  |||| | ||||
Db   1728 ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA 1787

Qy   1673 ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC 1732
               |  || |  | ||    ||||| |  ||||| | |||| |||   || | |||||||||
Db   1788 GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC 1847

Qy   1733 CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA 1792
             || | |||  |||||||       ||| || |||||||||| | |||  ||||||||   |
Db   1848 CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA 1907

Qy   1793 GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG 1852
             | |     |||||  | |  |     |||||  |  | || | |||||| |||||||||| |
Db   1908 GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG 1967
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Qy    1853 TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG 1912
           || ||  |||||  ||||||  | |      |     | ||  |||||||||||||||| |||| |
Db    1968 TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG 2027

Qy    1913 CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG 1972
           ||||  || || |   || |||||||||  || |  |   |||  |||  || |       |
Db    2028 CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC 2087

Qy    1973 TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG 2032
            | || | |||||| ||| |||||||  ||| |||||     | | || || || |  |||||||||||| |
Db    2088 GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG 2147

Qy    2033 AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC 2092
           ||||||  | ||||||||||||||||||||| |  | |||||| |  |  ||||| ||||||||||||| ||
Db    2148 AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG 2207

Qy    2093 AGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG 2152
           |||||||| |    ||||| |||  |  || |  |||||||      ||      |||||||
Db    2208 AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG 2267

Qy    2153 ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA 2212
           |||||||||||||||||||||||| ||| |||| ||  | | |||    |||  |||| ||| |
Db    2268 ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA 2327

Qy    2213 AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT 2272
           ||||  || |  | ||||||||||||||||||||||||| |||||||| |||| ||||| |||  |
Db    2328 AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG 2387

Qy    2273 ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA 2332
            | | |||||||| |||||||||||| |    |||| | | | || || |||||||
Db    2388 CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA 2447

Qy    2333 AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA 2392
           || |  || ||| |||| |||||  || || || |||| |||      |  |   ||  |
Db    2448 AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG 2507

Qy    2393 CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG 2449
           |  ||       | |  |    ||     ||     |   |    |||  |
Db    2508 CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA 2567

Qy    2450 CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA 2509
           ||| | |||| ||         ||||||||||||||| |   ||| || |||||||||    ||
Db    2568 CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA 2627

Qy    2510 GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC 2569
           |||| ||   |  |||     |||||     ||||||||| | || |||||| |||
Db    2628 GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA 2687

Qy    2570 TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC 2629
           ||||| |   | || | | ||||||||| |||| ||||||||| |  ||||||| |||
Db    2688 TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC 2747

Qy    2630 TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG 2689
           | |||||| || | |||    | |  ||||||| | ||||| |  ||  |||
Db    2748 TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG 2807
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Qy    2690 GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
                |||||  ||    ||           |       |    | || |   || ||||||
Db    2808 AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA 2860
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RESULT 15
US-10-218-779-3
; Sequence 3, Application US/10218779
; Publication No. US20040029222A1
; GENERAL INFORMATION:
; APPLICANT: Edinger, Shlomit
; APPLICANT: MacDougall, John
; APPLICANT: Millet, Isabelle
; APPLICANT: Ellerman, Karen
; APPLICANT: Stone, David
; APPLICANT: Gerlach, Valerie
; APPLICANT: Grosse, William
; APPLICANT: Alsobrook II, John
; APPLICANT: Lepley, Denise
; APPLICANT: Rieger, Daniel
; APPLICANT: Burgess, Catherine
; APPLICANT: Casman, Stacie
; APPLICANT: Spytek, Kimberly
; APPLICANT: Boldog, Ferenc
; APPLICANT: Li, Li
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Mishra, Vishnu
; APPLICANT: Patturajan, Meera
; APPLICANT: Shenoy, Suresh
; APPLICANT: Rastelli, Luca
; APPLICANT: Tchernev, Velizar
; APPLICANT: Vernet, Corine
; APPLICANT: Zerhusen, Bryan
; APPLICANT: Malyankar, Uriel
; APPLICANT: Guo, Xiaojia
; APPLICANT: Miller, Charles
; APPLICANT: Gangolli, Esha
; TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-214
; CURRENT APPLICATION NUMBER: US/10/218,779
; CURRENT FILING DATE: 2002-08-14
; PRIOR APPLICATION NUMBER: 60/253,834
; PRIOR FILING DATE: 2000-11-29
; PRIOR APPLICATION NUMBER: 60/250,-926
; PRIOR FILING DATE: 2000-11-30
; PRIOR APPLICATION NUMBER: 60/264,180
; PRIOR FILING DATE: 2001-01-25
; PRIOR APPLICATION NUMBER: 60/313,656
; PRIOR FILING DATE: 2001-08-20
; PRIOR APPLICATION NUMBER: 60/327,456
; PRIOR FILING DATE: 2001-10-05
; NUMBER OF SEQ ID NOS: 216
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
;   LENGTH: 2860
;   TYPE: DNA
;   ORGANISM: Homo sapiens

Query Match   34.0%;   Score 936.2;   DB 17;   Length 2860;
Best Local Similarity   61.7%;   Pred. No. 3.6e-230;
Matches 1662; Conservative   0;   Mismatches 938; Indels   93; Gaps   7;

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Qy      143 ACCCAGTGCCTGGTGCCAACCCGGACCTGCTTCCCCACTTCCTGGTGGAGCCCGAGGATG 202
            || |   | ||    ||     | || | ||| ||| |||||||| |||||| |||| |
Db      168 ACTCCTTCCCGTCAGCGCCAGCAGAGCCGCTGCCCTACTTCCTGCAGGAGCCACAGGACG 227

Qy      203 TGTACATCGTCAAGAACAAGCCAGTGCTGCTTGTGTGCAAGGCCGTGCCCGCCACGCAGA 262
            ||||| || |||||||||||| ||| ||| ||| |||    ||| | ||||||||| |||
Db      228 CCTACATTGTGAAGAACAAGCCTGTGGAGCTTCGCTGCCGCGCCTTCCCCGCCACACAGA 287

Qy      263 TCTTCTTCAAGTGCAACGGGGAGTGGGTGCGCCAGGTGGACCACGTGATCGAGCGCAGCA 322
            ||| |||||||||||||||| ||||||||| |||||     |||||||| |  ||   ||
Db      288 TCTACTTCAAGTGCAACGGCGAGTGGGTCAGCCAGAACGACCACGTCACACAGGAAGGCC 347

Qy      323 CAGACGGGAGCAGTGGGCTGCCCACCATGGAGGTCCGCATTAATGTCTCAAGGCAGCAGG 382
            || | | || || |||| |||||     ||| |||||  | | || || |||||||||||
Db      348 TGGATGAGGCCACCGGCCTGCGGGTGCGCGAGGTGCAGATCGAGGTGTCGCGGCAGCAGG 407

Qy      383 TCGAGAAGGTGTTCGGGCTGGAGGAATACTGGTGCCAGTGCGTGGCATGGAGCTCCTCGG 442
            | ||| || | | || ||||||||||||||||||||||||||||||||| |||||||| |
Db      408 TGGAGGAGCTCTTTGGGCTGGAGGATTACTGGTGCCAGTGCGTGGCCTGGAGCTCCGCAG 467

Qy      443 GCACCACCAAGAGTCAGAAGGCCTACATCCGCATAGCCAGATTGCGCAAGAACTTCGAGC 502
            |||||||||||||||   |||||| |||||||| |||   |||||||||||||||||| |
Db      468 GCACCACCAAGAGTCGCCGAGCCTACGTCCGCATCGCCTACCTGCGCAAGAACTTCGATC 527

Qy      503 AGGAGCCGCTGGCCAAGGAGGTGTCCCTGGAGCAGGGCATCGTGCTGCCCTGCCGTCCAC 562
            ||||||| |||| |||||||||||| |||||||||||| ||| |  |  | ||||| || |
Db      528 AGGAGCCTCTGGGCAAGGAGGTGCCCCTGGACCATGAGGTTCTCCTGCAGTGCCGCCCGC 587

Qy      563 CGGAGGGCATCCCTCCAGCCGAGGTGGAGTGGCTCCGGAACGAGGACCTGGTGGACCCGT 622
            |||||| |  | |||    ||||||||||||| ||||||   ||||||   |   |||||
Db      588 CGGAGGGGGTGCCTGTGGCCGAGGTGGAATGGCTCAAGAATGAGGATGTCATCGACCCCA 647

Qy      623 CCCTGGACCCCAATGTATACATCACGCGGGAGCACAGCCTGGTGGTGCGACAGGCCCGCC 682
            ||| |||| |||| |    |||| ||   ||||| |||   | | || |||||||||||||
Db      648 CCCAGGACACCAACTTCCTGCTCACCATCGACCACAACCTCATCATCCGCCAGGCCCGCC 707

Qy      683 TTGCTGACACGGCCAACTACACCTGCGTGGCCAAGAACATCGTGGCACGTCGCCGCAGCG 742
            |   | |||||| ||||| |||||||||||||||||||||||||||  ||| ||| |||
Db      708 TGTCGGACACTGCCAACTATACCTGCGTGGCCAAGAACATCGTGGCCAAACGCCGGAGCA 767

Qy      743 CCTCCGCTGCTGTCATCGTCTACGTGAACGGTGGGTGGTCGACGTGGACCGAGTGGTCCG 802
            || | || | |||||||||||||||||| || ||  ||||| |   ||| | |||||||
Db      768 CCACTGCCACCGTCATCGTCTACGTGAATGGCGGCTGGTCCAGCTGGGCAGAGTGGTCAC 827

Qy      803 TCTGCAGCGCCAGCTGTGGGCGCGGCTGGCAGAAACGGAGCCGGAGCTGCACCAACCCGG 862
            | |||| |   | | ||||||| || || ||||||||||| | ||||||||||||||| |
Db      828 CCTGCTCCAACCGCTGTGGCCGAGGCTGGCAGAAGCGCACCCGGACCTGCACCAACCCCG 887

Qy      863 CGCCTCTCAACGGGGGCGCTTTCTGTGAGGGGCAGAATGTCCAGAAAACAGCCTGCGCCA 922
            | || |||||||| || ||||| ||  ||||| ||| |       ||||||| || |||
Db      888 CTCCACTCAACGGAGGGGGCCTTCTGCGAGGGCCAGGCATTCCAGAAGACCGCCTGCACCA 947
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Qy     923 CCCTGTGCCCAGTAGACGGCAGCTGGAGCCCGTGGAGCAAGTGGTCGGCCTGTGGGCTGG 982
           || | |||||||| || ||     ||||    ||||||||||||||||| ||||| |    |
Db     948 CCATCTGCCCAGTCGATGGGGCGTGGACGGAGTGGAGCAAGTGGTCAGCCTGCAGCACTG 1007

Qy     983 ACTGCACCCACTGGCGGAGCCGTGAGTGCTCTGACCCAGCACCCCGCAACGGAGGGGAGG 1042
           | || ||||||||||| ||||| ||||||   | || ||||||  |||||||| |
Db    1008 AGTGTGCCCACTGGCGTAGCCGCGAGTGCATGGCGCCCCCACCCCAGAACGGAGGCCGTG 1067

Qy    1043 AGTGCCAGGGCACTGACCTGGACACCCGCAACTGTACCAGTGACCTCTGTGTACACAGTG 1102
           | ||| || || || || ||| | ||||| || || || || | ||    |
Db    1068 ACTGCAGCGGGACGCTGCTCGACTCTAAGAACTGCACAGATGGGCTGTGCATGCAACTGG 1127

Qy    1103 CTTCTGGCCCTGAGGACGTGGCCCTCTATGTGGGCCTCATCGCCGTGGCCGTCTGCCTGG 1162
           |        || | | |         | | | | | |||  |   ||| |    | |||
Db    1128 AGGCCTCAGGGGATGCGGCGCTGTATGCGGGGCTCGTGGTGGCCATCTTCGTGGTCGTGG 1187

Qy    1163 TCCTGCTGCTGCTTGTCCTCATCCTCGTTTATTGCCGGAAGAAGGAGGGGCTGGACTCAG 1222
           | ||| ||   ||    | | || || |||| ||     | | ||| |||
Db    1188 CAATCCTCATGGCGGTGGGGGTGGTGGTGTACCGCCGCAACTGCCGTGACTTCGACACAG 1247

Qy    1223 ATGTGGCTGACTCGTCCATT---CTCACCTCAGGCTTCCAGCCCGTCAGCATCAAGCCCA 1279
           | |  ||||||| ||  |     || ||     || ||||| ||||||| | | ||| |
Db    1248 ACATCACTGACTCATCTGCTGCCCTGACTGGTGGTTTCCACCCCGTCAACTTTAAGACGG 1307

Qy    1280 GCAAAGCAGACAACCCCCATCTGCT------CACCATCCAGCCGGACCTCAGCACCACCA 1333
           | |     ||||| || || ||     |  | | | || ||||| |   ||| |
Db    1308 CAAGGCCCAGTAACCCGCAGCTCCTACACCCCTCTGTGCCTCCTGACCTGACAGCCAGCG 1367

Qy    1334 CCACCACCTACCAGGGCAGTCTCTGTCCCCGGCAGGA----------------------- 1370
           || || |||||  ||     | | ||| ||||||
Db    1368 CCGGCATCTACCGCGGACCCGTGTATGCCCTGCAGGACTCCACCGACAAAATCCCCATGA 1427

Qy    1371 ----------------------TGGGCCCAGCCCCAAGTTCCAGCTCACCA--------- 1399
                                 ||||||| ||| || || || ||     ||
Db    1428 CCAACTCTCCTCTGCTGGACCCCCTTACCCAGCCTTAAGGTCAAGGTCTACAGCTCCAGCA 1487

Qy    1400 --ATGGGCACCTGCTCAGCCCCCTGGGTGGCGGCCGCCACAC----------ACTGCACC 1447
             | |||| | | || || ||     | | |||| | |          | ||
Db    1488 CCACGGGCTCTGGGCCAGGCCTGGCAGATGGGGCTGACCTGCTGGGGGTCTTGCCGCCTG 1547

Qy    1448 ACAGCTCTCCCACCTCTGAGGCCGAGGAGTTCGTCTCCCGCCTCTCCACCCAGAACTACT 1507
           || | || | |  ||| ||| |||  | || | |  | || |    |         | |
Db    1548 GCACATACCCTAGCGATTTCGCCCGGGACACCCACTTCCTGCACCTGCGCAGCGCCAGCC 1607

Qy    1508 TCCGCTC---------------CCTGCCCCGAGGCACCAGCAACATGACCTATGGGACCT 1552
           || | ||               |||||||||| | |  | | | ||   | || ||||
Db    1608 TCGGTTCCCAGCAGCTCTTGGGCCTGCCCCGAGACCCAGGGAGCAGCGTCAGCGGCACCT 1667

Qy    1553 TCAACTTCCTCGGGGGCCGGCTGATGATCCCTAATACAGGTATCAGCCTCCTCATCCCCC 1612
           |   || ||| || || ||  |||| | |||||     ||||| |||||| | ||  | |||
Db    1668 TTGGCTGCCTGGGTGGGAGGCTCAGCATCCCCGGCACAGGGGTCAGCTTGCTGGTGCCCA 1727

Qy    1613 CAGATGCCATACCCCGAGGGAAGATCTATGAGATCTACCTCACGCTGCACAAGCCGGAAG 1672
           | ||||| |||| || || | ||| |||| |||| |||| || |      | |||||| | |||
Db    1728 ATGGAGCCATTCCCCAGGGCAAGTTCTACGAGATGTATCTACTCATCAACAAGGCAGAAA 1787
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Qy    1673 ACGTGAGGTTGCCCCTAGCTGGCTGTCAGACCCTGCTGAGTCCCATCGTTAGCTGTGGAC 1732
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Db    1788 GTACCCTGCCGCTTTCAGAAGGGACCCAGACAGTATTGAGCCCCTCGGTGACCTGTGGAC 1847

Qy    1733 CCCCTGGCGTCCTGCTCACCCGGCCAGTCATCCTGGCTATGGACCACTGTGGGGAGCCCA 1792
                || | |||  |||||||   ||| ||  |||||||||   | |||  ||||||||  ||   ||
Db    1848 CCACAGGCCTCCTGCTGTGCCGCCCCGTCATCCTCACCATGCCCCACTGTGCCGAAGTCA 1907

Qy    1793 GCCCTGACAGCTGGAGCCTGCGCCTCAAAAAGCAGTCGTGCGAGGGCAGCTGGGAGGATG 1852
                | |       |||||  | | |   |||||| |   ||| |    | |||||  |||||||||| |
Db    1908 GTGCCCGTGACTGGATCTTTCAGCTCAAGACCCAGGCCCACCAGGGCCACTGGGAGGAGG 1967

Qy    1853 TGCTGCACCTGGGCGAGGAGGCGCCCTCCCACCTCTACTACTGCCAGCTGGAGGCCAGTG 1912
                || ||   |||||  ||||||  | |      |    | || |||||||||||||||||| |||| |
Db    1968 TGGTGACCCTGGATGAGGAGACCCTGAACACACCCTGCTACTGCCAGCTGGAGCCCAGGG 2027

Qy    1913 CCTGCTACGTCTTCACCGAGCAGCTGGGCCGCTTTGCCCTGGTGGGAGAGGCCCTCAGCG 1972
                ||||   || || |    ||  |||||||||| || |  | |   ||| ||| ||  |
Db    2028 CCTGTCACATCCTGCTGGACCAGCTGGGCACCTACGTGTTCACGGGCGAGTCCTATTCCC 2087

Qy    1973 TGGCTGCCGCCAAGCGCCTCAAGCTGCTTCTGTTTGCGCCGGTGGCCTGCACCTCCCTCG 2032
                 | ||| | |||||| ||| ||| |||||     | || || || |    |||||||||||| |
Db    2088 GCTCAGCAGTCAAGCGGCTCCAGCTGGCCGTCTTCGCCCCCGCCCTCTGCACCTCCCTGG 2147

Qy    2033 AGTACAACATCCGGGTCTACTGCCTGCATGACACCCACGATGCACTCAAGGAGGTGGTGC 2092
                ||||||  |  |||||||||||||||||||||||  |  |||||| |  |  |||||   ||||||||||  ||
Db    2148 AGTACAGCCTCCGGGTCTACTGCCTGGAGGACACGCCTGTAGCACTGAAGGAGGTGCTGG 2207

Qy    2093 AGCTGGAGAAGCAGCTGGGGGGACAGCTGATCCAGGAGCCACGGGTCCTGCACTTCAAGG 2152
                |||||||| |    ||||| |||  | ||  | |||||||       ||     |||||||
Db    2208 AGCTGGAGCGGACTCTGGGCGGATACTTGGTGGAGGAGCCGAAACCGCTAATGTTCAAGG 2267

Qy    2153 ACAGTTACCACAACCTGCGCCTATCCATCCACGATGTGCCCAGCTCCCTGTGGAAGAGTA 2212
                ||||||||||||||||||||||||  |||  |||| ||   | |||       ||| |||| ||| |
Db    2268 ACAGTTACCACAACCTGCGCCTCTCCCTCCATGACCTCCCCCATGCCCATTGGAGGAGCA 2327

Qy    2213 AGCTCCTTGTCAGCTACCAGGAGATCCCCTTTTATCACATCTGGAATGGCACGCAGCGGT 2272
                ||||  || | ||    |||||||||||||||||||| |||||||| ||||| |||||   |||  |
Db    2328 AGCTGCTGGCCAAATACCAGGAGATCCCCTTCTATCACATTTGGAGTGGCAGCCAGAAGG 2387

Qy    2273 ACTTGCACTGCACCTTCACCCTGGAGCGTGTCAGCCCCAGCACTAGTGACCTGGCCTGCA 2332
                 | | ||||||||| |||||||||||||  |   ||||       | |  || ||  ||||||
Db    2388 CCCTCCACTGCACTTTCACCCTGGAGAGGCACAGCTTGGCCTCCACAGAGCTCACCTGCA 2447

Qy    2333 AGCTGTGGGTGTGGCAGGTGGAGGGCGACGGGCAGAGCTTCAGCATCAACTTCAACATCA 2392
                  || | || ||| |||| ||||| ||||| || || || ||||   |||       | | ·  ||   |
Db    2448 AGATCTGCGTGCGGCAAGTGGAAGGGGAGGGCCAGATATTCCAGCTGCATACCACTCTGG 2507

Qy    2393 CCAAG---GACACAAGGTTTGCTGAGCTGCTGGCTCTGGAGAGTGAAGCGGGGGTCCCAG 2449
                |  ||       |  | |     ||    ||        |      ||      ||| |
Db    2508 CAGAGACACCTGCTGGCTCCCTGGACACTCTCTGCTCTGCCCCTGGCAGCACTGTCACCA 2567

Qy    2450 CCCTGGTGGGCCCCAGTGCCTTCAAGATCCCCTTCCTCATTCGGCAGAAGATAATTTCCA 2509
                ||| | |||| ||       |||||||||||||||||||||||  |  |||  || |||||||||       ||
Db    2568 CCCAGCTGGGACCTTATGCCTTCAAGATCCCACTGTCCATCCGCCAGAAGATATGCAACA 2627

Qy    2510 GCCTGGACCCACCCTGTAGGCGGGGTGCCGACTGGCGGACTCTGGCCCAGAAACTCCACC 2569
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              |||| ||  | |||        |||||      |||||||||||   |  || |||||| |||
Db    2628 GCCTAGATGCCCCCAACTCACGGGGCAATGACTGGCGGATGTTAGCACAGAAGCTCTCTA 2687

Qy    2570 TGGACAGCCATCTCAGCTTCTTTGCCTCCAAGCCCAGCCCCACAGCCATGATCCTCAACC 2629
              ||||| |   | || |   | |||||||| ||||  | |||||||| |    ||||||| |||
Db    2688 TGGACCGGTACCTGAATTACTTTGCCACCAAAGCGAGCCCCACGGGTGTGATCCTGGACC 2747

Qy    2630 TGTGGGAGGCGCGGCACTTCCCCAACGGCAACCTCAGCCAGCTGGCTGCAGCAGTGGCTG 2689
              |  ||||| || | |||        | | ||  ||||||  |    |||||      ||  |||   |
Db    2748 TCTGGGAAGCTCTGCAGCAGGACGATGGGGACCTCAACAGCCTGGCGAGTGCCTTGGAGG 2807

Qy    2690 GACTGGGCCAGCCAGACGCTGGCCTCTTCACAGTGTCGGAGGCTGAGTGCTGA 2742
              ||||| ||   ||        |        |   | || | || ||||||
Db    2808 AGATGGGCAAGAGTGAGATGCTGGTGGCTGTGGCCACCGACGGGGACTGCTGA 2860


Search completed: March  6, 2005, 17:03:48
Job time : 1506.37 secs
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